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Command line parameters:

MODEL=frame+_p2n.model -DEV-xlp

MODEL=frame+_p2n.model -DEV-xlp

-Q-/Ggn2_1/USPTO_Spool_/US09827854_runat_11032003_101610_27486/app_query.fasta_1.3576

-DB-GenEmb1 -QFMT-fastap -SUFFIX=p2n.rge -MIRMATCH=0.1 -LOOPCL=0 -LOOPEXT=0

-UNITS-bits -STARF=1 -END=-1 -MATREIX=blosum62 -TRANS=human40.cdi -LIST=45

-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL

-OUTFMT=pto -NORM-ext -HEAPSIZE=500 -MINILEN=0 -MAXLEN=200000000

-USEER-US09827854_eCGN_1_1_13008_erunat_11032003_101610_27486 -NCPU=6 -ICPU=3

-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120

-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7

-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Maximum Match 100%
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29: em_vi:*
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36: em_htg_vrt:*
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37: em_htg_vrt:*
38: em_sy:*
39: em_htgo_hum:*
40: em_htgo_mus:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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                                                                               ThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla
                                  ArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                           LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGlnGlnLeu
                                                                                                               CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
                       CGGCTGGGCGGGACATGGAGGACGTGCGCGGCGGCGCGTGCAGTACCGCGGCGAGGTG
                                                                    ACCCCGGTGGCGGAGGAGGCGGGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
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OS Homo sapiens (human)
PN JP 2001017028-A/3
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128919
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                                                     PATENT: JP 1985118189-A 1 2
MITSUBISHI CHEM IND LTD
OS human
PN JP 1985118189-A/1
PD 25-JUN-1985
PF 29-NOV-1983 JP 1983224
PI TERANISHI YUTAKA, TAKA
PI TERANISHI YUTAKA, TAKA
KIMURA MASAKO,
PI IKEDA YASUKO
PC C12N15/00,C07H21/04//C
CC strandedness: Double;
CC topology: Linear;
CC hypothetical: No;
CC anti-sense: No;
CC anti-sense: No;
CC anti-sense: No;
CC anti-sense: No;
CC TOS TISSUE_type=1
FH Key 15.90
FT CDS JFT CDS
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 1110)
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DNA FRAGMENT
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JP 1985118189-A/1.
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coding human apolipoprotein
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C12N15/00,C07H21/04//C12P21/00;
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29-NOV-1983 JP 1983224980
/product='apolipoprotein
                                                                                           tissue_type=liver;
    Location/Qualifiers
                                                 15.
                         69.
                                              /product='apolipoprotein
.68
                        /product='apolipoprotein E3
.965
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TERANISHI YUTAKA, MATSUI YASUSHI, IKEDA YASUKO, KIMURA MASAKO C12P21/00, A61K35/74, A61K37/04, C12N15/00, (C12P21/00, C12R1:19),
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Patent: JP 2001017028-A; 2 23-JAN-2001;
Patent: JP 2001017028-A; 2 23-JAN-2001;
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/2
PD J2-JAN-2001
PF 28-APR-2000 JP 2000128919
PR 21-INOBU FUJITA, HIROKI HAMANAKA, YUKC
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AOLK67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC
C12N15/09//C07K14/775, PC
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Breslow, J.L.
Synthesis, intra
apolipoprotein
                                                                        2 (bases 250 to 777) Wallis, S.C., Rogne, S., Gill, L., Markham, A., Edge, M., Woods, D. Williamson, R. and Humphries, S.
The isolation of cDNA clones for human apolipoprotein E and detection of apoE RNA in hepatic and extra-hepatic tissues EMBO J. 2 (12), 2369-2373 (1983)
                                                                                                                                                      J. Biol. Chem. 83082756
                                                                                                                                                                                   Breslow, J.L., \ McPherson, J., \ Nussbaum, A.L., \ Williams, H.W., \ Lofquist-Kahl, F., \ Karathanasis, S.K. \ and \ Zannis, V.I. \ Identification \ and \ DNA \ sequence \ of \ a \ human \ apolipoprotein
                                                                                                                                                                                                                                                                       apolipoprotein; apolipoprotein low density lipoprotein.
                                Zannis, V.I., McPherson, J.,
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1 (bases 355 to 1156)
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Karathanasis, S.K.

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Woods, D.,

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GTGCAGGCTGCCGTGGGCACCAGCGCCCCCCTGTGCCCCAGCGACAATCAC
                   ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
                                                                                 AGCTGGTTCGAGCCCCTGGTGGAAGACATGCAGCGCCAGTGGGCCGGGCTGGTGGAGAAG
                                                                                                     SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                                                                                                                                     CTGGAGGAGCAGGCCAGCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCCCGCCTCAAG
                                                                                                                                                                                         LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
                                                                                                                                                                                                                                                      AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCCAAG
                                                                                                                                                                                                                                                                            SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Apo E is a component of normal human very low density lipoproteinemia. There are six human apo E phenotypes known to result from a single structural gene, three of the common alleles being epsilon-4, epsilon-3 and epsilon-2. This sequence appears to be of the epsilon-3 allele. [1] argues that the apo E polymorphism involves mutations in the structural coding region; for example the epsilon-2 phenotype which is characterized by hyperlipoproteinemia is thought to result from a c to t change (arg to cys) at base 586 below [3], [5]. The sequence shown is 57% homologous with human apo A-I and 81% homologous with rat apo E. For the epsilon-4 sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       [1] corrected in [J. Biol. Chem. 258, 11422-11422 (1983)].
Biol. Chem. 258, 11422-11422 (1983)] correction of [1]. [4]
epsilon-3 and variant.
[5] epsilon-2 allele.
[6] sites; mutation-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 (sites)
Rall,S.C. Jr., Newhouse,Y.M., Clarke,H.R., Weisgraber,K.H., McCarthy,B.J., Mahley,R.W. and Bersot,T.P.
Mye III hyperlipoproteinemia associated with apolipoprotein bhenotype E3/3. Structure and genetics of an apolipoprotein E
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84185684
                                                                                                                                                                                                                                                                                                                                                                                            Apo E is located on chromosome 19 --Jackson, Bruns and Breslow, PNAS USA 81, 2945-2949 (1984) -- and is believed to be linked to the Apo C-II gene (see separate entry).

The two mutations causing type III hyperlipoproteinemia (apolipoprotein E phenotype E3/3) produces substitutions of Arg for Cys at amino acid 112 and Cys for Arg at amino acid position 142. Draft entry and printed copy of sequence for [1] kindly provided by L.L. Gill, University of Aberdeen.
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Humphries, S.E.,
                                                                                                                                                                                                                                                                                                                          Complete source information: Human liver [1],[J. Biol. Chem. 258, [4] and blood [5], cDNA to mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        [J. Biol. Chem. 258, 11422-11422 (1983)] is too extensively revise relative to [1] to record the revisions in the FEATURES table. The sequence below is that of [J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       see
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Isolation and characterisation of a variant allele of the gene for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J. Biol. Chem. 259 (10), 6498-6504 (1984) 84212473
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/product="preapolipoprotein
/protein_id="AAB59546.1"
                                                                                                                                                      /tissue_type="liver and blood"
1. .1156
                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                /codon_start=1
                                                                          /gene="APOE"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hisiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy,
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Submitted (20-FBB-2001) National Institutes of Health, Mammalian Gene Collection (MCC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
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                                           LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
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MOTIMOTO, H. and Teranishi, Y.
METHOD FOR PRODUCING APOLIPOPROTEIN
PATENT: JP 194315392-A 1 15-NOV-1994;
MITSUBLISHI KASEI CORP
OS HOMO SAPIENS (human)
PN JP 194315392-A/1
PD 15-NOV-1994
PF 11-JUN-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF 120V-1994
PC C12P21/02,CO7K13/00,C12N5/10,C12N15/
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PC (C12N15/10,C12R1:91);
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PC (Strandedness: Double;
PC topology: Linear;
FH Key Location/Qualifiers
FH Source /organism-'Homo sapi
FT sig_peptide 15. 68
FT mat_peptide 69. 968
FT mat_peptide 69. 968
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                                                                                   LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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E08423.1 GI:217654
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Homo sapiens.
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MORIMOTO HIRONORI, TERANISHI YUTAKA
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/organism="Homo sapiens"
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                                                                                    Fujita,S., Hamanaka,H., Fu
Apo E humanized mammal
Patent: JP 2001017028-A 1
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/1
PD 23-JAN-2001 JP 200012
PF 28-APR-2000 JP 200012
PR SHINOBU FUJITA,HIROKI
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                                                        PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE AOLIK67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC C12N15/09/C07K14/775, PC (C12N5/10, C12R1:91), C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
                                                                                                                                                                                               Mammalia;
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mmalia; Eutheria; Primates;
(bases 1 to 1156)
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                 Location/Qualifiers
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Location/Qualifiers
  /organism="Homo
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                                                                                                                                                                          Fukui, Y.
                                                        C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
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Avalon Pharmaceuticals (US)
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Mammalia; Eutheria;
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                                                                                                                    Similarity:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/odb_xref="taxon:9606"
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                                                                                               SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
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115975
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        1 (bases 1 to 1157)
Attie, A.D., Gretch, D.G., Sturley, S.
Production of recombinant proteins
Patent: US 5472858-A 1 05-DEC-1995;
Location/Qualifiers
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                                                                                             GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                         CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAAGTCACCCAAGAACTGAGGGCG
                                      LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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/translation="MKYLWAALLVTFLAGCQAKVEQPVEPETEPELRQQAEGQSGQPW
/translation="MKYLWAALLVTFLAGCQAKVEQPVEPETEPELRQQAEGQSGQPW
ELALGREWDYLRWVQTLSEQVQEELLSPQVTQELTTLMDETMKELKAYKSELEEQLSP
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KLRKRLLRDADDLQKRLAVYQAGAREGAERGVSAIREBLGPLVEQGRVBAATVCSLAS
CLUCBBAQALGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQISLQAEAFQ
                                                                                                                                                                                                                                                                                                                                                                /product="proapolipoprotein
/protein_id="CAA32092.1"
/db_xref="GI:38055"
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/clone_lib="lambda g
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        Human apolipoprotein E (epsilon-4 allele) gene, complete cds. M10065 J03053 J03054 M10065.1 G1:178852 M10065.1 G1:178852 Alu repeat; allelic variation; apolipoprotein; apolipoprotein lipoprotein; repeat region; very low density lipoprotein. Human DNA [2], [1].
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sapiens
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(epsilon-4
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A potential TATA box is found at positions 1014-1018, and a potential polyadenylation signal at 4616-4621.

[2] and [1] had slight differences in the boundary positions for the Alu repeats and their flanks; the boundary positions indicated in [1] have been used in the FEATURES table below. Draft entries and clean copies were kindly supplied by J.M. Taylor, Gladstone Laboratories, San Francisco, and by J.P. Levine, Rockefeller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Emi,M., Wu,L.L., Robertson,M.A., Myers,R.L., Hegele,R.A.,
Williams,R.R., White,R. and Lalouel,J.M.
Genortyping and sequence analysis of apolipoprotein E isoforms
Genomics 3 (4), 373-379 (1988)
89212602
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85216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 5515)
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J. Biol. Chem. 260 (10), 6240-6247 (1985)
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.Emi, 19-AUG-1988.
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                                /citation=[2]
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1789. .1793
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202. .203
                                                                                                                                                                                                                                                                                 /citation=[2]
/replace=""
                                                                                                                                                                                                                                                                                                                                            /note="direct repeat
963. .967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    complement(300. .345)
/note="direct repeat flanking"//
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/replace=""
                                                                                                                     /note≂"apo E mRNA intron 1 [2],[1]"
1362. .1363
                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Alu repeat
                                                                                                                                                                                                                                                                                                                                                                                                                                                            complement(346. .635)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                            /note="apo E mRNA [2],[1]"
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.680)
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/replace=""
2887. .2893
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/replace=""
3007. .3199
                                                                                        4342
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/number=2
1914. 3006
/note="g in epsilon-4; a in epsilon-3 [2]"
4708. .4709
/citation=[2]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="g in epsilon-4; a 3200. .3780
                                                                                                                                                                          /note="gac
[3]"
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3191
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2527. .2886
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complement(2/3)
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complement(2105. .2429)
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1871. .1913
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1871. .1913
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ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH"
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VAEETRARLSKELQAAQARLGADMEDYRGRLYQYRGEYQAMLGQSTEELRYRLASHLR
KLRKRLLRDADDLQKRLAYYQAGAREGAERGLSAIRERLGPLYEQGRYRAATYGSLAG
                                                                                                                  /note="c in epsilon-4;
                                                                                                                                                                                                                                                                                                                     3817.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      /citation=[2]
/replace=""
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3568. .3569
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/replace=""
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2947. .2948
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join(3018. .3199,3781. .4495)
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/protein_id="AAB59397.1"
/db_xref="GI:178853"
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join(1871. .1913<sub>.</sub>3007. .3199,3781. .4498)
                                                                                                                                                                                                                                                                                                                                            'number=3
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                                                                                                                                                                                                                                                                                                                                                                       note="preapolipoprotein
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eat flanking Alu repeat 5'
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78			78	Qy
3688	rcredecerececerrecrrerececrerreger	TGCATCTGTCTGTCTCCTTCT	3629	Db
78			78	Qy
3628	CTTCTTCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTC	ACCTTGCCCGGCCTCCTAGCTCC	3569	Дb
78			78	Qy
3568	CTCGGCCTCCCAAAGTGCTGGGATTAGAGGCATGAGC	CTGGGCTCAAGCGATCCTCCCGC	3509	DЪ
78			78	Qy
3508	rggggtctcactgtgttgcccaggctggtcttgaactt	CTTTTATATAGAGACAGAGAGATG	3449	Дb
78			78	Qy
3448	CACACTCGTCCTGGCTCTGTCTCTGTCCTTCCCTAGCT	CTTCCCTTCTGACTCAGTCTCTC	3389	Db
78			78	Qy
3388	3CTCTCTGGAATTCTCTCTCTCAGCTTTGTCTCTCTCT	CCATTTCTGACTCCTGGCTTTAGC	3329	밁
78			78	Qy
3328	SGGGGCCTGGGTCTCTGCTGGTTCTAGCTTCCTCTTC	TCTGCCCCTGTCGCTAAGTCTTG	3269	В
78			78	Qy
3268	CTGGTGGGCGGCTATACCTCCCCAGGTCCAGGTTTCAT	CCCATCCTGGCCCTTGACCCTCC	3209	рь
78			78	Qγ
3208	TCCCAGGTCACCCAGG	GAGCAGGTGCAGGAGGAGCTGCTCAGC	3150	DЪ
78	SerGln	. H	63	Qy
62 3149	rgPheTrpAspTyrLeuArgTrpValGlnThrLeuSer 	ArgTrpGluLeuAlaLeuGlyArgPheT 	43 3090	ОУ
42 3089	ProGiuLeuArgGlnGlnThrGluTrpGlnSerGlyGln 	AlaValGluThrGluProGluPr 	23 3030	Qу
3029	ThrPheLeuAlaGlyCysGlnAlaLysValGluGln 	AAGGACCCTGACCCGACCTTGAACT	2970	D dq
•				
10 2969	euVal         TOTGCCCCTAGGTAGCTAGATGCCTGGACGGGGTCAG	LysValLeuTrpAlaAlaLeuLe	2 2910	Qу
	DE4 (1-5515)	-854-14 (1-317) х нимаро	09-827	US-
	Length: 5515  Matches: 307  Conservative: 1  Mismatches: 8  Indels: 214  Gaps: 2	1.24e-66 1.27e-66 1.27e-66 1.27e-66 1.27e-66 1.27e-66 1.27e-66 1.24e-66 1.38e-7.08 1.24e-66 1.38e-7.08 1.24e-66 1.38e-7.08 1.24e-66 1.38e-7.00	gnmentd. No. re: cent St Loca	Ali Pre Sco Per Bes Que DB:
[1]	614768)   Grant flanking Alu repeat 3' copy 695048)   peat [1]   peat flanking Alu repeat 5' copy .100 g 1206 t	at_region complement /note="direction complement /note="ali at_region complement /note="direction complement /note	repe repe repe	BAS
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Fgapop 6.0 , 1
Delop 6.0 , 1
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1: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1980.
2: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1981.
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## SUMMARIES

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## ALIGNMENTS

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                                  CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
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GENAISSANCE PHARM INC

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The patent discloses novel genetic variants of human apolipoprotein CC E (APOE) gene. The invention also relates to compositions and methods CC for haplotyping and/or genotyping the APOE gene. The haplotyping CC methods of the invention are useful for improving the efficacy and CC reliability of several steps in the discovery and development of CC drigs for treating diseases associated with APOE activity, e.g. CC familial dysbetalipoproteinaemia, type III hyperlipoproteinaemia, CC atherosclerosis, and Alzheimer's disease. They are useful to validate PAPOE as a candidate agent for treating a specific condition or disease condicted to be associated with APOE activity and in the design of CC clinical trials of candidate drugs for treating a specific condition CC or haplotyping methods are useful to screen for compounds targeting CC APOE to treat a specific condition or disease associated with APOE activity. The present sequence is a CDNA encoding human APOE protein.

CC APOE gene is located on chromosome 19q13.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleoti pair at specific polymorphic sites for two copies of gene -
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                 apolipoprotein-E (or who produce abnormal forms of this molecule) and therefore are likely to suffer from hyperlipidemia, resulting arterlosclerosis. It can also be used to raise antisera for detecting the protein deficiency or production of abnormal forms.
                                                                                    New DNA sequence coding expression vectors and t
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RESULT 5
ABA83113
The invention relates to methods for diagnosing and prognosing ovarian CC tumours in an individual via the detection and measurement of the CC expression of ovarian tumour marker genes (ABA83181-ABA83182, ABA83180, CC ABA83182 and ABA83184) or segments thereof (ABA83181-ABA83122, ABA83179, CC ABA83181 and ABA83184) or segments thereof (ABA83181-ABA83129, ABA83189). The methods of the invention are useful for CC detecting an ovarian tumour in a patient, for identifying an individual CC at increased risk for developing ovarian cancer, in prognostic tests for CC monitoring a patient in remission from ovarian cancer and in tests for CC monitoring disease status in a patient being treated for ovarian cancer. CC The methods can additionally be used to identify a particular tumour as being an ovarian tumour (i.e., an epithelial ovarian tumour selected from CC serous cystadenoma, borderline serous tumour, serous cystadenoma, concertinoma, cystadenoma, borderline mucinous tumour, mucinous cystadenoma, borderline mucinous tumour, mucinous cystadenoma, borderline mucinous tumour, mucinous cystadenoma, cystadenofibroma, adenofibroma and Brenner CC tumour. The ovarian analysis of gene expression) and were found to be coverexpressed in a broad variety of ovarian epithelial tumour cells coverexpressed in a broad variety of ovarian pathers denominated to proliferation and in protein folding, and many of these are membrane-CC implicated in immune response pathways, in the regulation of cell cocalised or secreted. In addition to their use as diagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    serous cystadenocarcinoma; mucinous cystadenocarcinoma; mucinous cystadenoma; borderline mucinous tumour; endometrioid carcinoma undifferentiated carcinoma; clear cell adenocarcinoma; cystadenofibroma; adenofibroma; Brenner tumour; serial analysis of gene expression; SAGE; immune response pathway; cell proliferation regulation; protein folding; membrane localised; secreted; therapeutic target; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene
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prognostic markers, the ovarian tumour marker genes or their encoded proteins may be used as therapeutic targets for the treatment and prevention of ovarian cancer. Sequences ABA83081-ABA83122, ABA83180,

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                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to an ApoE hum present sequence is the coding sequence for in the method of the present invention. The can be used for screening for agents useful Alzheimer's disease and arteriosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for treating arteriosclerosis -
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DB; AAB80997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       humanized mammalian cell useful for screening or treating or preventing Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 13-14; 22pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  coding
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for human ApoE3, which was us
The ApoE humanised mammalian
ful for treating or preventin
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RESULT 7
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hypertriglyceridaemia; low density lipoprotein; LDL; ds.
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                                                                                                                                                                                                                                                                                                         GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                                                                                                                                                      AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCCAAG
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/*tag= a
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                         Location/Qualifi 61..1014
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                                                                                                                                                                                                                                                                                                                         The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atheroselerosis, treating or regressing atheroselerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LI receptor or is at risk for developing atheroselerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
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04-OCT-2000;
05-APR-2001;
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(UYBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                     New apolipoprotein E polypeptide and nucleic acid, cholesterol, delaying the onset of or treating ath mammal, without inducing hypertriglyceridemia -
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GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                                                                                                                                ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
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DB; AAE13294.
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RESULT 8
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hypertriglyceridaemia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New apolipoprotein E polypeptide and nucleic accholesterol, delaying the onset of or treating mammal, without inducing hypertriglyceridemia
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P-PSDB;
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05-APR-2001; 2001US-0827854.
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                             CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
                                                                    CGGCTGGGCGGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
                                                                              ArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgGlyGluVal
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The present invention relates to an ApoE humanised mammalian cell. The present sequence is the coding sequence for human ApoE2, which was us in the method of the present invention. The ApoE humanised mammalian can be used for screening for agents useful for treating or prevential Alzheimer's disease and arteriosclerosis.
                                                                                    Disclosure;
                                                                                                                New apoE humanized ruseful for treating arteriosclerosis -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             New apolipoprotein E polypeptide and nucleic ac cholesterol, delaying the onset of or treating mammal, without inducing hypertriglyceridemia
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04-OCT-2000; 2000US-0679088.
05-APR-2001; 2001US-0827854.
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                                                                                                                                                                                                                                                                   SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                                                                                                                                                                                                                                                                                         GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAGCAAGCGGTGGAGACAGAGCCCGAGCCGAGCTGCGCAGACCGAGTGGCAGAGC
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                                                                                 standard;
(first
                                                                                 DNA;
  entry)
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900 280 840

260

780

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720

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540

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160 480 140 420 120 100

360

300 80 240 60 180 40

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Percent Similarity:
Best Local Similarity
Query Match:
DB:
                                                                                                                                                                                                                                                                                      Alignment
Pred. No.:
Score:
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                                                                                                                                                                                                                                                       Local Similarity:
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04-OCT-2000;
05-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is a human apolipoprotein E (apoE) allele, apoE2* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (L receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or haddens to the content of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (KOSP-)
(UYBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New apolipoprotein E polypeptide and nucleic acid, cholesterol, delaying the onset of or treating ath mammal, without inducing hypertriglyceridemia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zannis
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                                                                                                                                                                                                                                                                                                           No.:
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                                                                                                        GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                     GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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B; AAE13297.
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; 2000US-0679088.
; 2001US-0827854.
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Matches:
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RESULT 12
AAT06957
ID AAT06
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                                                  Recombinant; human; apolipoprotein-E; ApoE; insect cells; Manduca sexta; Autographica californica nuclear polyhedros haemolymph; lipid complex; biologically active; ds.
                                                                                          Human apolipoprotein-E (ApoE)
                                                                                                                                                         AAT06957
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                                                                                                                                                         standard;
 Location/Qualifiers 62..119
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                                                    polyhedrosis virus; ds.
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                                                                                                                                                                                                                                                                       Percent Similarity:
                                                                                                                                                                                                                                                                                                                                             Recombinant human apolipoprotein-E (ApoE) (AAR86791) can be produced by preparing a genetic construct (contg. an ApoE-encoding sequence, e.g. AAT06957, and flanking regulatory sequences enabling the protein to be expressed in insect cells), which is then introduced into a Manduca sexta larva (using a recombinant Autographica californica nuclear polyhedrosis virus) and recovering the protein from the haemolymph of the larval host. The ApoE produced is in a form sufficiently complexed with lipids to be biologically active, which cannot be achieved in insect cell cultures, and can therefore be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-JUN-1991;
                                                                                                                                                                                                                                                                                                                      Sequence 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Manduca
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                                                                                                                                                                                                                                                                                           No.:
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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGlnGlnLeu
                                                                       MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
                                     CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
                                                                                                                                                                                      ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
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                                                                                                                                                                                                                                                              Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          of recombinant apo:lipoprotein E in insects a sexta larvae with recombinant Autographica r polyhedrosis baculovirus vector.
                                                                                                                                                                                                                                                                                                                                      therpeutic applications
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CC Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO care for the printed and the control of the printed careful control of the printed careful carefu
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progression of benign prostatic hyperplasia (BPH), or screening (II) for or identifying an agent that modulates the onset or progression of BPH. The method is based on changes in gene expression in BPH tissue isolated from patients exhibiting different clinical states of prostate hyperplasia as compared to normal prostate tissue. (I) comprises detecting the expression levels of one or more genes in prostate cells from the subject that are differentially regulated compared to normal prostate cells. (II) comprises preparing a first gene expression profile of BPH cells or BPH-like cell population, exposing the cells to the agent, preparing a second gene expression profile of the agent exposed cells, and comparing the first and second gene expression profiles.

(I) is useful for diagnosing the onset or progression of BPH. (II) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENE-)
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05-JUN-2001;
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                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                                patient
                                                                                                                                                                                                                                                                                                                                                  Identifying drugs for and diagnosing benign prostatic hyperplasia, detecting expression levels of one or more genes in prostate cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            useful for identifying an agent that modulates the onset or progression of BPH. The methods are useful to present information identifying the expression level in a tissue or cells, by comparing the expression level of genes given in the specification in the tissue or cells to the level of expression of gene in the database, and displaying the expression levels of at least one gene in the tissue or cell sample compared to the expression level in BPH. Agents using (II) are useful for treating BPH or prostate cancer. ABK64106-ABK64860 represent human benign prostatic hyperplasia gene sequences of the invention.
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gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;
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-Q-ggn2_1/USPTO_Spool_US09877854_runat_11032003_101611_27510/app_query.fasta_1.3576
-OB=ISSUED_FATENTS_NA -OPMT=fastap | SUFFIX=p2n.rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALICM=200 -THR_SCORE=pèt -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09827854 @CGN_1_1_224_@runat_11032003_101611_27510 -NCPU=6 -ICPU=3
-NO_XIPXY -NO_MAMP -LARGEQUERY -NEGL_SCORES=0 -WAIST -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPDEXT=0.5 -FGAPDP=6 -FGAPEXT=7
-YGAPDP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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18 - 07 - 849 - 389 - 6

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Compugen Ltd.
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                              APPLICATION NUMBER: US/07/709
FILING DATE: 19910604
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 9629
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
INFORMATION FOR
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: Quarles &
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ZIP: 53701-2113
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Sturley, Stephen I
Beckage, Nancy E
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US-09-397-766-24
US-09-397-681-24
US-09-495-444-24
US-08-952-796-1
US-07-959-946-4
US-08-952-796-1
US-08-968-751-3
US-08-968-751-3
US-08-952-796-1
US-08-952-796-1
US-08-952-796-1
US-08-952-796-1
US-09-183-861-7
US-09-183-861-7
US-09-183-861-34
US-08-952-765-7
US-09-183-861-34
US-08-195-487-3
US-08-195-487-3
US-08-195-27-126
US-08-195-27-126
US-08-195-27-126
US-08-195-27-126
US-08-353-105-2
US-08-961-537-126
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SEQUENCE CHARACTERISTICS: LENGTH: 1157 base pairs

STRANDEDNESS: double

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DB:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PUBLICATION INFORMATION:

MCLean, J W
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JOURNAL: Journal of Biological Chemistry
VOLUME: 259
PAGES: 6498-6504
DATE: 1984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION:
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                                                                                                                    ArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                                              ThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla
                                LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
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                                                                                                          CGGCTGGGCGGCGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
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                     AAGCTGCGTAAGCGGCTCCTCCGCGATCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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Mismatches:
Indels:
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                                                                       REFERENCE/DOCKET NUMBER: TA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEPAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                              APPLICATION NUMBER: US 60/027, 338
FILING DATE: 11-CCT-1996
PRIOR APPLICATION NUMBER: US 60/046,094
                                                                                                                                                                                                                                                                        FILING DATE: Concurre
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
                                                                                                                                                                              FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Piedrahita, Jorge A APPLICANT: Bazer, Fuller W TITLE OF INVENTION: Composition TITLE OF INVENTION: Generation
                                                            SEQUENCE CHARACTERISTICS:
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TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: line
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                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/949,155
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STREET: P.O. BOX 4433
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 CTCCTCAAAGGCTGGTTCGAGCCTCTGGTTGGAAGACATACGGCGCCAGTTGGGCCGGGCTG
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          ArgLeuLysSerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeu
                                                                                                                                  GGCCAGCCGCTGCGCGAGCGCGCGAAGCCTGGGGCCAGAAGCTGCGCGGACGGCTGGAG
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Query Match:
DB:
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US-09-819-964-5
; Sequence 5, Application US/09819964
; Patent No. 6369294
; GENERAL INFORMATION:
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                                                                                                                                          US-09-827-854-14 (1-317) x US-09-819-964-5 (1-1126)
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                                                                                                                                                                                                                                Percent Similarity:
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ETILING DATE: <Unknown>
APPLICATION NUMBER: US 60/046,094
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD,
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Bazer, Fuller W
                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 418-3000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Compositions and Methods
                               GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu 37
                                                                                        LENGTH: 1126 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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CITY: Houston
STATE: TX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/819,964 FILING DATE: 28-Mar-2001
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70.31%
72.49%
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                Sequence 51, Application US/08949155 Patent No. 6271436 GENERAL INFORMATION:
                                                APPLICANT: Piedrahita, Jorge A
APPLICANT: Bazer, Fuller W
TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: Generation of Transgenic
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
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CITY: H
COUNTRY: US
ZIP: 77210-4433
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                   STATE:
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TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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78
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                                                                                      TrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrp
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                          TTGGGGGGAAAAAACTTTTTCTTTTTTCTTTCATTTGACTTCATGTCTTGCTTTCCTTTCC
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Matches:
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US-09-813
; Sequence 51, Application
; Patent No. 6369294
; GENERAL INFORMATION:
APPLICANT: Piedrahita, Jorge A
Bazer, Fuller W
Bazer, Fuller W
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Bazer Fuller W
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US-09-819-964-51
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            NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE
STREET: P.O. BOX 4433
                                                                     TITLE OF INVENTION: Compositions
Generation of
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CTGAGTACCAGGGCCGGCCAGCCGCTGCGCGAGCGCGGAAGCCTGGGGCCAGAAGCTG
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                                                                                                                                                                                                                                                                                             GlnTrpAlaGlyLeuValGluLysValGlnAlaAlaValGly---ThrSerAlaAlaPro
                                                                                                                                                                                                                                                                                                                                                      GluAlaPheGlnAlaArgLeuLysSerTrpPheGluProLeuValGluAspMetGlnArg
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CITY: Houston
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Query Match:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51:
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LENGTH: 4267 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION: NAME: Hibler, David W.
                                                                                                                                                                                                                                ValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGlu
                                                                                                                                                                                                                                                                                          TrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                           GAGCCGGGG------CCGCCGCCGGAGGTGCACGTGTGGTGGAGGAGCCCAAG
                                                                                                                                                                                                                                                                                                                                                                 GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu
                                                                                                                                                                                                                                                                                                                                                                                               ATGGCTGTGAACTGGCTCGGT--------GCAGGATGCCGGACAGAGGAC
                                                                                                  CCCTCCTGGCGAACCGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 6 FILING DATE: 09-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/949,155 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: <Unknown>
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Conservative:
Mismatches:
Indels:
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RESULT 6
US-08-726-306A-28
           Sequence 28, Application US/08726306A
Patent No. 5958684
GENERAL INFORMATION:
APPLICANT: van Leeuwen, Frederik Willem
APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
TITLE OF INVENTION: DIAGNOSIS METHOD AND REAGENTS
NUMBER OF SEQUENCES: 189
   NUMBER OF SEQUI
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 96,0.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9110
TELEPAS: (617) 345-9111
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: GB 95/20080.4
FILING DATE: 02-Oct-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
FILING DATE: 01-Jan-1996
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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 422
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STREET: 1 Fin
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02111
                                                                                                                                                                                                                                               182
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 660 base pairs
LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                                                                                                                                                                                                                         GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                          ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120
                                                                                                                                      LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                                                                                                                                                                                                                                                                             GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
                                                             ACCCCGGTAGCGGAGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC
                                                                                                                          CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAAGAACAACTG
                                                                                                                                                                                    CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
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Matches:
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                                                                               Query Match:
DB:
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US-07-849-389-6
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                                                                                                         Best Local Similarity:
                                                                                                                   Percent Similarity:
                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 1678
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEPAX: (703)883-4109
TELEX: 899149
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6,
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                                                                                                                                               . No . :
                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                    MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: HORNES, Erik APPLICANT: UHLEN, Mathias
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: USE FILING DATE: 19920519 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
SOFTWARE: Patentl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE:
COMPUTER: II
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ZIP: 22313-0299
                              87
                                                                                                                                                                                                              NAME/KEY:
                                                                                                                                                                                                                                      ORGANISM:
                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                         STRANDEDNESS:
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STATE: Virginia
                                                                                                                                                                                                                                                                                                                 LENGTH:
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GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAGCTGCGTAAGCGGCTCCTCCGCGATCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                           linear
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98.18%
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         Query Match:
                       Best Local Similarity:
                                Percent Similarity:
                                            Score:
                                                       Pred. No.:
                                                                   Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                          APPLICATION NUMBER: 08/406,19
FILING DATE: MATCh 17, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth A.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: SQI-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
TENERRATION FOR SEO IN NO. 24.
                                                                                                                                                                  TELEFAX: (617)227-594
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL
                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                                                                                                                                                 CLASSIFICATION: PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kvster, Hubert TITLE OF INVENTION: DNA D
                                                                                                    MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61
                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: March
                                                                                                              TOPOLOGY:
                                                                                                                                     TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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                                                                                                                                   nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                            60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                        LAHIVE & COCKFIELD
                                                                                                                                                                                                                                                                                                                                                                        Floppy disk
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                                                                                                                      single
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18, 1996
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         Length:
Matches:
Conservative:
Mismatches:
Indels:
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US-09-827-854-14 (1-317) x US-08-617-256-24 (1-252)

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              Alignment Scores: Pred. No.:
                                                             US-09-287-141-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 08/617,256
                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: K ster, TITLE OF INVENTION:
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CORRESPONDENCE ADDRESS:
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                                                                          TOPOLOGY: 1
DLECULE TYPE:
                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002D
                                                                                                                                                                                                                                                                                                NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/6 FILING DATE: 18-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: SOFTWARE: Patent
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                                                                                                                      nucleic acid
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                                                                                                                                                                                   (617)227-7400
(617)227-5941
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DB:
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                                             TELEFAX: 619-587-53b0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
570 hase pairs
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GENERAL INFORMATION:
                                                                                                                                       REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002G
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
                                                                                                                                                                                                              APPLICATION NUMBER: 08/406, FILING DATE: 03-MAR-1995 ATTORNEY/AGENT INFORMATION: NAME: Seidman, Stephanie L
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APPLICATION NUMBER:
FILING DATE: 18-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
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                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                STRANDEDNESS:
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Best Local Similarity:
Query Match:
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GENERAL INFORMATION:
TELEFAX: 858-587-5360
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                               APPLICATION NUMBER: 08/617
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                         NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 247
TELECOMMUNICATION INFORMATION:
TELEPHONE: 858-450-8400
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatit
OPERATING SYSTEM: DOS
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TITLE OF INVENTION: DNA Diagnostics
NUMBER OF SEQUENCES: 33
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CURRENT APPLICATION DATA:
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CLASSIFICATION:
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; MOLECULE TYPE:
US-09-504-245-24
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US-09-287-682-24
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Best Local Similarity:
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                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/4(
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
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No.:
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APPLICATION NUMBER: 08/6
FILING DATE: 18-MAR-1996
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MEDIUM TYPE: Diskette
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TELECOMMUNICATION INFORMATION:
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                         NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
SOFTWARE: Patentl
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ZIP: 92037-9103
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STATE: CA
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/287,682
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SYSTEM: DOS
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Heller Ehrman White & McAuliffe
Heller Expansion White & McAuliffe
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INFORMATION FOR SEQ ID NO: 24
SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                        Sequence 24, Application US/09287679
Patent NO. 6258538
GENERAL INFORMATION:
APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08
APPLICATION NUMBER: 08
FILING DATE: 18-MAR-18
                                                                            SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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TELEFAX: 619-587-5360
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                     COUNTRY: USA
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                 ADDRESSEE: Hell
STREET: 4250 Exe
CITY: La Jolla
STATE: CA
COUNTRY: USA
2IP: 92037-9103
                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
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STRANDEDNESS: si
TOPOLOGY: linear
                                                                                                                        OPERATING SYSTEM:
                                                                                                                                          COMPUTER:
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(77) TD NO: 24:
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                                                                            US/09/287,679
                 08/617,256
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Gaps:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                      GENERAL INFORMATION:
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TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24
                                                      COUNTRY: USA
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                        APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA Diagnostics NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/4
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
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             COMPUTER: IBM CON OPERATING SYSTEM: SOFTWARE: Patentl
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TELEFAX: 619-587-5360
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               PatentIn Release #1.0, Version #1.25
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                           IBM Compatible SYSTEM: DOS
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Indels:
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Query Match:
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Best Local Similarity:
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                                                                                                Sequence 24, Application US/09287681
Patent NO. 6277573
GENERAL INFORMATION:
APPLICANT: K Ster, Hubert
TITLE OF INVENTION: DNA Diagnostics Based
NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELECHIONE: (617)227-7400
TELECAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White
STREET: 4250 Executive Square,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
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TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/40
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                 149
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002I
              COUNTRY:
                            STATE:
                                          CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 GCACGGCTGTCCAAGGAGCTGCAGGCGGGCGCAGGCCCGGCTGGGCGCGGACATGGAGGAC
                                                                                                                                                                                                                                                                                                                   AspAlaAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla 188
                                                                                                                                                                                                                                                                                                                                                                                GluGluLeuArgValArgLeuAlaSerHisLeuArgLysLeuArgLysArgLeuLeuArg 168
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                       La Jolla
E: CA
92037-9103
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                                                         & McAuliffe
7th Floor
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Matches:
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                      Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/617,256
FILING DATE: 18 MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
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                                                   181
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                          189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 619-450-8400 TELEFAX: 619-587-5360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette
AspAlaAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla 188
                                                 GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGGCCCGCGAGGGCCCC
                                                                                                                                                                                                                                                                                                                                                                                             TYPE:
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n Release #1.0, Version
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Matches:
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Search completed: March 14, 2003, 20:18:22 Job time : 40.7343 secs

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-Q=/cgn2_1/USPTO_spool/US09827854/runat_11032003_101612_27578/app_query.fasta_1.3576
-Q=/cgn2_1/USPTO_spool/US09827854/runat_11032003_101612_27578/app_query.fasta_1.3576
-DB=Published_applications_NA -QFWT=fastap -SUFFIX=p2n.rnpb -MINMATCH=0.1
-LOOPEXY=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -QUTFWT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=2000000000 -USER=US09827854 | GCGN_1_1_298_derunat_11032003_101612_27578
-NCPU=6 -ICPU=3 -NO_XLPXY -NO_MMAP | LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
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-MODEL=frame+_p2n.model
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9 US-09-870-759-129

9 US-09-802-640-17

10 US-09-827-854-8
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Sequence 7, Appli
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454, Appl 12, Appl 12, Appl 11, Appl 11, Appl 760, App 1760, Appl 1760, Appl 10, Appl 10, Appl 13, Appl 133, Appl 134, Appl 5420, Ap 4726, Ap 4726, Ap 9395, Ap 9395, Ap
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                                                                          CTGGAGGAGCAGGCCCAGCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCCCGCCTCAAG
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Patent No. US20020177551A1

GENERAL INFORMATION:
APPLICANT: TERMAN, David S
TITLE OF INVENTION: COMPOSITIONS AND METHODS
FILE REFERENCE: 870759
CURRENT FILING DATE: 2002-01-14
PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR REPLICATION NUMBER: US 60/208,128
PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR PRILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 166
SOFTWARE: Patentin version 3.1
SEQ ID NO 129
LENGTH: 1156
TYPE: DNA
ORGANISM: Homo Sapiens
FEATURE:
NAME/KEY: CDS
NAME/KEY: CDS
OTHER INFORMATION:
US-09-870-759-129
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 17
LENGTH: 1156
TYPE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: GENES AND POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISEASE AND THEIR USE
FILE REFERENCE: 24736-2048
CURRENT APPLICATION NUMBER: US/09/802,640
CURRENT FILING DATE: 2001-03-09
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bonsal Aruna APPLICANT: Kleyn Patrick
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TYPE: DNA
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LOCATION: (61)...(1014)
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Condesterol levels with File Reference: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriak
                                        SEQ ID NO 8
                                                                                                                                                                                               Sequence 8, Application US/09827854 Patent No. US20020123093A1
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TYPE: DI
TYPE: DNA
ORGANISM: Homo
09-827-854-8
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; OTHER INFORMATION: I
; NAME/KEY: unsure
; LOCATION: 46
; COTHER INFORMATION: a
US-10-044-090-454
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TITLE OF INVENTION: GENES DIFFERENTIALLY EXF
FILE REFERENCE: PA-0028 US
CURRENT APPLICATION NUMBER: US/10/044,090
CURRENT FILING DATE: 2002-01-09
NUMBER OF SEQ ID NOS: 850
SOFTWARE: PERL Program
SEQ ID NO 454
LENGTH: 1291
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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US-10-044-090-454
; Sequence 454, Application i
; Patent No. US20020137081A1
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cohlesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US/09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. US20020123093A1
GENERAL INFORMATION:
APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriako
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LENGTH: 1156
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APPLICANT: Zannis, vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: ChoLesterol levels without inducing hyp
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                                         Sequence 9, Application Patent No. US20020123093 GENERAL INFORMATION:
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                                                                      SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
                                                                                                   GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILLING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 1156
TYPE: DNA
ORGANISM: Homo sapiens
US-09-827-854-11
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APPLICANT: Zannis, Vassilis
APPLICANT: Kyprecs, Kyriakos E.
APPLICANT: Kyprecs, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without induci
FILE REFERENCE: 07180,/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
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0	; ORGANISM: Homo sapiens US-09-954-456-760 Alignment Scores:	Pate 760 1157	PRIOR FILING DATE: 2000-09-27 PRIOR APPLICATION NUMBER: US/60/235,863 PRIOR FILING DATE: 2000-09-27 NUMBER OF SEQ ID NOS: 2276	FILING DATE: 2000-09-27 APPLICATION NUMBER: US/60/235, FILING DATE: 2000-09-27 APPLICATION NUMBER: US/60/235,		FILING DATE: 2000-09-25 APPLICATION NUMBER: US/60/235, FILING DATE: 2000-09-25 APPLICATION NUMBER: US/60/235,	FILING DATE: 2000-09-18 APPLICATION NUMBER: US/60/234, FILING DATE: 2000-09-20 APPLICATION NUMBER: US/60/234,	REFERENCE: 68929 NT APPLICATION N NT FILING DATE: APPLICATION NUM	GENERAL INFORMATION: APPLICANT: Young, Paul APPLICANT: YOUNG, Paul TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Canc TITLE OF INVENTION: Sets	RESULT 9 US-09-954-456-760 ; Sequence 760, Application US/09954456 ; Patent No. US20020115057A1	Qy 301 ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 317	Qy 281 SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300 	Qy 261 LeuGluGluGlnAlaGInGInIleArgLeuGInAlaGluAlaPheGInAlaArgLeuLys 280	Qy 241 SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys 260	Qy 221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240	Qy 201 ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220	601 CAGGCCGGGGCCCGAGGGCGCCCGAGCGCCTCAGGGGCATCCGCGAGCGCCTGGGG 660
RESULT 10 US-09-880-107-2244 US-09-880-107-2244; Application US/09880107 ; Sequence 2244, Application US/09880107 ; Patent No. US20020142981A1 ; GENERAL INFORMATION:	Qy 301 ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 317	Qy 281 SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300	Qy 261 LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 280	Oy 241 SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys 260	Qy 221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240	Qy 201 ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220	Qy 181 GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200	Qy 161 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr 180	Qy 141 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160	Qy 121 ArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgGlyGluVal 140	Qy 101 ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120	Qy 81 LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu 100	Qy 61 LeuSerGluGlnValGlnGluCeuLeuSerSerGlnValThrGlnGluLeuArgAla 80	Qy 41 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 60	Qy 21 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40	Qy 1 MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20	US-09-827-854-14 (1-317) x US-09-954-456-760 (1-1157)

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APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-W0
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT PILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-01-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2244
LENGTH: 1157
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ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10
LENGTH: 1156
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                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Coholesterol levels without inducing hypertriglyceridemia
                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 07180/004003
                                                                                                                                                                                                                                                                                                            LENGTH: 11
TYPE: DNA
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ment Sim	JILT 12 39-967-01 39-967-01 39-967-01 39-967-01 39-967-01	301 V	281 S     901 A	261 L     841 C	241 SI 1 781 AI	221 L 1 721 C	201 P: 661 C	181 G   	161 L     541 A	141 G   	121 Ar   -   421 CG	101 T 361 A	81 L
cores: ilarity	1)13-5 5, Appl 0, US200 1)10 Stan TIN STAN TIN STAN 1 INVENT 1 INVENT	ValGlnAl          GTGCAGGC	SerTrpPh        AGCTGGTT	euGluG]         !TGGAGG!	SerArgTl         AGCCGGAC	euGlnG]          TACAGG!	ProLeuVa           CCCTGGT	lnAlaG1          AGGCCGG	LysLeuAr         AAGCTGCG	GlnAlaMetLe             CAGGCCATGCT	rgLeuGl         GCTGGG	ThrProVa	LeuMeta:          TGATGG:
••	ESULT 12  \$7-99-967-013-5  Sequence 5, Application US/09 Patent No. US20020045840A1  GENERAL INFORMATION: METHOD F  TITLE OF INVENTION: MUMBER: CORRENT APPLICATION NUMBER: COURTENT APPLICATION NUMBER: 60  PRIOR APPLICATION NUMBER: 60  PRIOR FILING DATE: 2000-05-2  NUMBER OF SEQ ID NOS: 91  SOFTWARE: FastSEQ for Window  SEQ ID NO 5  LENGTH: 41907  TYPE: DAM  ORGANISM: Homo sapiens  S-09-967-013-5	LaAlaVa          TGCCGT	rpPheGluProI               GTTCGAGCCCC	uGluGluGlnAlaGlnG             GGAGGAGCAGGCCCAGC	gThrargas             GACCCGCGA	luArgAl         \GCGGGC	oLeuValGluGlnGlyA              CCTGGTGGAACAGGGCC	66 – Y	TA I	00 - M	yAlaA        CGCGG	ω L	AspGluThrMetLysGluE 
4.61e- 1370.5 60.00%	nn US/09 140A1 140A1 1Fr. Vinc ETHOD F AND USE 1-022001 IUMBER: 2000-15-0200 00-05-291 Window	lGlyTh        GGGCAC	OLeuVal	aGlnGl         CCAGCA	o v	aGlnAl         CCAGGC	nGlyAr        .GGGCCG	laArgGluGlyAlaGl	ysArgLeuLeuArgAs            	lyGlnSe         ACCAGAG	PMetG1         CATGGA	.uGluTh        GGAGAC	nrMetLy         CATGAA
113 0	96701 ent p OR GE S THE S THE J206, 3	lnAlaAlaValGlyThrSerAla	GluA       GAAG	nIleAr         GATACG	uAspGl	LeuGlnGluArgAlaGlnAlaTrpGlyGluAr 	gValAre	yAlaGli          CGCCGAC	uArgAsi         ccgcgAi	rThreli         cacceac	SPMetGluAspValA	AlaGluGluThrArgAlaA 	sGluLe
Leng Matc Cons	EOF 2967,C	AlaProVal	spMetGlnArgGlr              ACATGCAGCGCCAC	InIleArgLeuGlnAlaGluAlaPheG 	ArgLeuAspGluValLysGluGlnVal. 		ArgValArgAlaAlaThrValGly	GluArgGlyLeuSerA              GĀGCGCGGCCTCAGCG	SPAlaAspAspLeuGlnLys	lnSerThrGluGluLeuAr 	LArgGlyAr          TGCGGCCG	O H	euLysAlaTyrLys 
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41 30 re: 0	9 OF AP	SerAspA         !AGCGACA	lnTrpAlac          \GTGGGCCG	laPheG	/alAlac         TGGCGG	gAlaArgMetGlu             CGCGCGGATGGAO	lyserLeu          GCTCCCTG	Laile	BlnLysA        AGAAGT	rgValArgLeuAlas             GGTGCGCCTCGCCT	rgLeuValGlnTy           CCTGGTGCAGT <i>F</i>	SluLeud          AGCTGC	SerGluI         CGGAAC
907	OE DNA	AsnHis          AATCAC	aGlyLeuVai           :GGGCTGGT	lnAla/       AGGCC	lAlaGluValArg <i>I</i>              GCGGAGGTGCGCC	fetGluG        \TGGAGG	euAlaGlyG          !TGGCCGGCC	ArgGluAr          CGCGAGCG	ArgLeuA        TGCCTGG	laserHisL	rArgo 	lnAla <i>i</i>            AGGCGG	LysSerGluLeuGluGl               AAATCGGAACTGGAGGA
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GCAGGCCGGCTGGGCCGGACATGGAGGACGTGYGCGCCGCCTGGTGCAGTACCGCGG	GlnAlaArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgG	uGlnLeuThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAl	-ArgAlaLeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGl 	CCCCATCCCAGCCCTTCTCCCCGCCTCCCACTGTGCGACACCCTCCCGCCCTCTCGGCCG		GCCTCTGCCCCGTTCCTCTCTCCCCTCTTGGGTCTCTCTGGCCTCATCCCCCATCTCGCCCG		TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTCTGTCTCTCTC		$\tt CCTCCCAAAGTGCTGGGATTAGAGGCATGAGCCACCTTGCCCGGCCTCCTAGCTCCTTCT$		$\tt CTCACTGTGTTGCCCCAGGCTGGTCTTGAACTTCTGGGCTCAAGCGATCCTCCCGCCTCGG$		CGTCCTGGCTCTGTCCTGTCCCTAGCTCTTTTATATAGAGACAGAGAGATGGGGT		${\tt TGGAATTCTCTCTCAGCTTTGTCTCTCTCTCTTCTCTTC$		$\tt CCTGGGTCTCTGCTGGTTCTAGCTTCCTCTTTCCCATTTCTGACTCCTGGCTTTAGCTCTC$		GGCGGCTATACCTCCCCAGGTCCAGGTTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGGG			SerGlnValThrGlnGluLeu	TrpAspTyrLeuArgTrpValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSer 	LeuArgGlnGlnThrGluTrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPhe 	PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProGlu	-854-14 (1-317) x US-09-967-013-5 (1-41907)	1 Similarity: 60.00% . Mismatches: 6 ch: 86.69% Indels: 194 caps: 1
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LENGTH: 786
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                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Nucleic Acids, Proteins FILE REFERENCE: PA104
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                                                                                                                                                                                                                                SOFTWARE: PatentIn version SEQ ID NO 374 LENGTH: 478
                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign TITLE OF INVENTION: Sets FILE REFERENCE: 689290-73
                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/964,824A CURRENT FILING DATE: 2001-09-27
                                                                                                                                                          NAME/KEY: misc_feature LOCATION: (1)...(478) OTHER INFORMATION: n=a.
                                                                                                                                                                                                ORGANISM: Homo sapiens FEATURE:
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RESULT 15
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Sequence 2491, Application U;
Patent No. US20020142981A1
GENERAL INFORMATION:
APPLICANT: HOTHE, Darci T.
APPLICANT: Vockley, Joseph
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc
                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:

APPLICANT: Horne, Darci T.

APPLICANT: Vockley, Joseph G.

APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-W0
CURRENT APPLICATION UMMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR APPLICATION NUMBER: US 60/337,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PATEENTIN VET. 2.1
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-UNITS-bits -STARP=1 -END=-1 -MAYRIX=bilosum62 -TRANS-human40.cd1 -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTPMT=pto -NORM-ext -HEAPSIZE-500 |-MINLEN=10000000000
-USER-USO9827854_CGN_1_1_8826_crundt_11032003_101610_27495 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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## ALIGNMENTS

REFERENCE AUTHORS TITLE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BI670367.	RESULT 1
<pre>1 (bases 1 to 1027) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens	human.	EST.	B1670367.1 GI:15584600	B1670367	mRNA sequence.	603292738F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5312024 5',	BI670367 1027 bp mRNA linear EST 12-SEP-2001		

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Contact: Robert St
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Email: cgapbs-r@mail.
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/lab_host="DH10B"
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/clone="IMAGE:5312024"
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                                                                                                                                                                                                               Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium ()
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                         DNA sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLCM1718 row: b column: 10
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/clone_lib="NIH_MGC_49"
/tissue_type="melanotic melanoma, |
/lab_host="DH10B (phage-resistant)
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                                                                                                            ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
                                                                                                                                                                                                  GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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                                                            LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMet-Gl
                                                                                                                                                                                  CAGGCCGGGGCCCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
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                                            AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
                                                                                                                                                                                                                                                                                                                      CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
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                                                                                                                      GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                   GAGCAAGCGGTGGAGACAGAGCCGGAGCCGAGCTGCGAGCAGACCGAGTGGCAGAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11798 row: a column: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein
Toshiyuki and Plero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consorti
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NIH-MGC http://mgc.nci.nih.gov/.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: Salt-xhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched
                                                                                                                                                                                                                                                                                                                                                                                        for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library." 284 c 359 g 119 t
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:5314844"
/clone_lib="NIH_MGC_96"
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/db_xref="taxon:9606"
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                                                                                                                                                                            mRNA sequence.
BG761746
BG761746.1 GI
EST.
         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTP/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium ()
DNA Sequencing by: Incyte Genomics, Inc.
                                                                                       1 (bases 1 to 938)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                       BG761746
602717942F1 NIH_MGC_49
                                                                   Contact: Robert Strausberg, Ph.D.
                                                                               Unpublished (1999)
                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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CAGGCCGGGGCCCGAGGGCGCGAGCGCGTGCTCAGCGCCATCCGCGAGCGCCTGGGG
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http://image.lln1.gov
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/Clone_lib="NIH_MGC_49"
/clone_lib="NIH_MGC_49"
/tissue_type="melanotic melanoma, high MDR (cell line)"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: xhor; Site_2:
/note="Organ: skin; Vector: pOTB7; Site_1: xhor; Cloned into EcoRI/Xhor sites using the following 5; adaptor:
GCACGAG(G). Size-selected >500bp for average insert size
1.8xb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
II RT (Life Technologies). Note: this is a NIH_MGC
Library. | "
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11762 row: a
column: 11
High quality sequence stop: 762.
Location/Qualifiers
1 0042
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National Institutes of Health, Mammalian
Unpublished (1999)
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1 (bases 1 to 942)
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                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5301010"
/clone_lib="NIHLMGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
                                                                                                                                                                                                      /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-Xhor (gtcgag pBluescript KS+); Site_1: Fire_7: Fire_
                                                                                                                for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

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mRNA sequence.
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BM042094.1 GI:
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ACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
                                                                                                                                  CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAGGTCACCCAGGAACTGAGGGCG
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Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1875 row: i column: 03 High quality sequence stop: 792.
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human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, |
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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/lab_host="DHIOB (phage-resistant)"
/note="organ: skin; Vector: poTBP; Site_1: XhoI; Site_2:
_note="organ: skin; Vector: potential organisms of colored into EcoRI/XhoI sites using the following 5 adaptor:
_gcCACGAN(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NHH_MGC Library."

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/db_xref="taxon:9606"
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BG472299
BG472299.1 GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                       found through the I.M.A.G.E. Consortium/LLNL Plate: LLCM1419 row: k column: 08 High quality sequence stop: 848.
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National Institutes of Health,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                 170
             /clone_lib="NiH_MGC_16"
/tlssue_type="retinoblastoma"
/tlssue_type="retinoblastoma"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2: Cloned into EccRI/XhoI sites using the following 5/
adaptor: GGCAGG(G). Library constructed by Ling Hong
in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-CDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
                 b
                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4645759"
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                                             CCGGCCTCAAGAGCTGGTTCGGAGCCCTGGGTGGAAGACATG
                                                                                        CCGCAAGCTGGAGGAAGCAGGCCAGCAGATACGGCTGCAGGCCGAGGGGCCTTCCAGGGC
                                                                                                    gAlaLysLeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAla---PheGlnAl
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EST.
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Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Miklos Palkovits, M.D., Ph.D.

cDNA Library Preparation: Michael J. Brownstein (NHGRI)

Toshiyuki and Piero Carninci (RTKEN)

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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http://image.llnl.gov
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Location/Qualifiers
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a 273 c 359 g 114 t
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/clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DHIOB"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                   CAAGCTGAAGAAACAGCCCAGAAATACGCCTTA---GCCGAAGGCTTCAGG
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602649975F1 NIH_MGC_40 H
mRNA sequence.
BG774871
BG774871.1 GI:14045188
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                                                                            cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1612 row: e column: 20 High quality sequence stop: 813.
                                                                                                                                                                                          Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
I (bases 1 to 817)
III-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
Unpublished (1999)
                                                                                                                                                            Tissue Procurement: DCTD/DTP
                                                                                                                                                                       Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                 human.
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4760827"
/clone_lib="NIH_MGC_40"
/tissue_type="carcinoma, c
/lab_host="DH10B (phage-re
                                                            Location/Qualifiers
1. .817
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Homo sapiens
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ArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgGlyGluVal

140

459 160 120

399

100

80 279

339

CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG

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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu

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698 240

758

ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro

CCCCTGGTGGAACAGGGCCGCGTGCGGGCCTGCACTGTGGGCTCCCTGGCCGGCAG-CCG

579

519 180

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Best Local Similarity:
Query Match:
                                                                                                                                                      US-09-827-854-14 (1-317) x BG774871
                                                                                                                                                                                                                    Percent Similarity:
                                                   100
                       41
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                                                 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                                                   Scores:
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                                                                                                                                                                                                                                                                                           /note="Organ: prostate; Vector: pOTB7; Site_1: XhoI; Site_2: ECORI; CDNA made by Oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCAGCAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
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1178.50
96.93%
96.17%
74.54%
                                                                                                                                                      (1-817)
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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251
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AUTHORS
TITLE
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GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                               Email: cgapbs r@mail.nih.gov

Tissue Procurement: Miklos Palkovits, M.D., Ph.D.

CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAMAI1726 row: g column: 10

High quality sequence stop: 783.
Location/Qualifiers
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 811)
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EST.
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603244936F1 NIH_MGC_96
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National Institutes of Health, Mammalian
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1177.00
97.97%
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REFERENCE
AUTHORS
TITLE
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                                                                                                                                                 found through the I.M.A.G.E. Consortium/LINL http://image.llnl.gov plate: LICM1790 row: g column: 17 High quality sequence stop: 829.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 845)

NIH-MCC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   602763768F1 NIH_MGC_42 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GI:14177059
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cDNA clone IMAGE:4899112
         pOTB7; Site_1: XhoI;
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Query Match:
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                                                                                                                                                                       GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                                                                                                                                                          LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                                                                                                                                                                                                                                                               LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
            SerArgThr-ArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
                                                   CTACAGGAGCGGCCAGGCTG---GGGCGAGCGGTGCGCGCGGATGGAGGAGATGGGC
                                                                                                     CCCCTGGTGGAACAGGGCCGCGTGCGGGCCGC-ACTGTGGGGCTCCCTTGGCGGCAG-CCG
                                                                                                                                                           CAGGCCGGGGCCCGAGGGCCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
                                                                                                                                                                                                             AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAGGTCACCCAGGAACTGAGGGCG
AGCCGGACCGCGA----CGCCTGGACGAGGTGAAGGAGCAGGTGCCGGAGGTGCGCCCAA
                                                                                                                                                                                                                                                                                                                                                                                                                       ACCCCGGTGGCGGAGGAGACGCGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
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                                           sLeuGluGluGln
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
CDNA Library Preparation: Michael J.
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.
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BG707147
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National Institutes of Health, Mammalian
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1140.00
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Brownstein
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                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
BI551475
BI551475.1 GI
                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consorti
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603194314F1 NIH_MGC_95
                                                                                                            DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11691 row: 1 column: 04
                                                                                                                                                                                                                                         Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
                                                                                                                                                                                                                                                                                              Mammalia; Eutheria; Primates; 1 (bases 1 to 919)
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                                                                                    quality sequence stop: |
Location/Qualifiers
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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В Qy ρς γΩ

Qy 밁 QΥ

В

172 a /note="Organ: brain; Vector: pBluescriptR (modified pBluescript RS+); Site\_1: BamHI; Site\_2: SalI-XhOI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.5 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Captrapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMM/NHGRI, National Institutes of Health). Note: this is a NIH\_MGC Library." a 270 c 363 g 114 t

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Percent Similarity:
Best Local Similarity:
Query Match:
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CAGCCGGACCCGGGACG - - CTGGACGAAGTGAAGAGCACGTGGGCGGAGGG - TGCGCCAA
                                       GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                                                                                                                                                                 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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          ySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
                                                                                                                                 CAGGCCGGGGGCCGATGGCGCCGAGCGCGTGCTCAGCGCCATCCGCGAGCGCCTGGGG
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93.26%
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BM728696
 Percent Similarity:
Best Local Similarity:
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TITLE
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                                                     No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BM728696 706 bp mRNA linear EST 01 UI-E-EJO-aiu-1-08-0-UI.r1 UI-E-EJO Homo sapiens cDNA clone UI-E-EJO-aiu-1-08-0-UI 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Program for Rat Gene Discovery and Mapping
University of Iowa
451 Eckstein Medical Research Building Iow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genetics (www.resgen.com).
Seg primer: M13 Reverse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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319 335 9565
                                                                                                                         123
                                                                                                                 /note="Organ: eye; Vector: pT73-Pac (Pharmacla) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; UI-E-EJO is a subtracted cDNA library constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was lighted to an EcoR I adaptor, digested with Not I, and cloned directionally into pT7T3-Pac vector. The oligonuclectide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tags for this library are: fetal eyes, AGAATCAAGA; lens, CGATTAAGTG; retina, CCGCG; Retina Foveal and Macular, GTCC; RPE and Chorold, ACCTA. This library was created for the program, Gene Discovery in the Visual System, supported by National Eye Institute (NEI)."
                                                                                                                         þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="UI-E-EJ0-aiu-1-08-0-UI"
/clone_lib="UI-E-EJ0"
/tissue_type="fetal eyes, lens, eye anterior segment,
optic nerve, retina, Retina Foveal and Macular, RPE a
Choroid"
                                                                                                                                                                                                                                                                                                                                                                                                                                /dev_stage="fetal and adult"
/lab_host="DH10B (Life Technologies) (Tl phage resistant)"
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/db_xref="taxon:9606"
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98.73%
98.73%
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                                                 REFERENCE
AUTHORS
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                                                                       Carninci, P.
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US-09-827-854-14 (1-317) x BM728696 (1-706)
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                                                                                                                                                 AKO10261.1 GI:12845571
HTC; CAP trapper.
Mus musculus (strain:C57BL/6J) ES cells cDNA to mrNA,
Clone_lib;RIKEN full-length enriched mouse cDNA library
                                                                                                                                                                                                                                             AK010261 1110 bp mRNA linear HTC 19-JAN-2002
Mus musculus ES cells cDNA, RIKEN full-length enriched library,
clone:2400003C12:apolipoprotein E, full insert sequence.
High-efficiency full-length cDNA
Meth. Enzymol. 303, 19-44 (1999)
                                                                                                                                     clone:2400003C12.
                                                                             Eutheria;
                                                                                             Metazoa;
                                     and Hayashizaki, Y
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                                                                                                                                                                                                           Submitted (10-JUL-2000) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Gemomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama Kanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.go.jp, URL:http://genome.gsc.riken.go.jp/, Tel:81-45-503-9222,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Marchionni, L., Mashima, J., Mazzarelli, J., Mombaerts, P., Nordone, P., Ring, B., Ringwald, M., Rodriguez, I., Sakamoto, N., Sasaki, H., Sato, K., Schonbach, C., Séya, T., Shibata, Y., Storch, K.F., Suzuki, H., Toyo-oka, K., Wang, K.H., Weitz, C., Whittaker, C., Wilming, L., Wynshaw-Boris, A., Yoshida, K., Hasegawa, Y., Kawaji, H., Kohtsuki, S. and Hayashizaki, Y.
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                                              cDNA library was prepared and sequenced in Mouse Genome 
Encyclopedia Project of Genome Exploration Research Grou
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                                                                                                                                                                  Please visit our web site (http://genome.gsc.riken.go.jp/)
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                                                                                                                                      further details.
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                                                                                                                                    ArgLeuGlyAlaAspMetGluAspValArgGlyArgLeuValGlnTyrArgGlyGluVal 140
                                                                                                                                                                               GTCCAGTGGCGGAGAGACACGGGCCAGGCTGGGCAAAGAGGTGCAGGCGGCACAGGCC
                                                                                                                                                                                                                  ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
                                                                                                                                                                                                                                                                                                                                 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                                                 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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	ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsn 316	301	Qy
942	GGCTGGTTCGAGCCAATAGTGGAAGACATGCATCGCCAGTGGGCAAACCTGATGGAAAAG	883	Db
300		281	Qy
882	ATGGAGGAACAGACCCAGCAAATACGCCTGCAGGCGGAGATCTTCCAGGCCCGCCTCAAG	823	DЪ
280	LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys	261	Qy
822	AACCAGGCCCGTGACCGCCTAGAGGAGGTGCGTGAGCACATGGAGCAGGTGCGCTCCAAG	763	Db
260		241	Qy
762	ATGTGCGATCGCGCCCAGGCTTTTGGTGACCGCATCCGAGGGCGGCTGGAGCAAGTGGGC	703	DЬ
240		221	Qy
702	CCTCTGGTGGAGCAAGGTCGCCAGCGCACTGCCAACCTAGGCGCTAGGGCCGCCAGCCT	643	Db
220	ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro	201	Qy
642	AAGGCAGGGCACGCTAGGGCGCCGAGCGCGGTGTGAGTGCCATCCGTGAGCGCCTGGGG	583	Db
200	GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly	181	Qy
582	AAGATGCGCAAGCGCTTGATGCGGGATGCCGAGGATCTGCAGAAGCGCCTAGCTGTGTAC	523	Db

Search completed: March 14, 2003, 20:13:47 Job time: 1296.69 secs

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Command line parameters:

-MODEL-frame+_p2n.model -DEV=xlp
-Q-cgn2_1/USPTO_spool/US09827854/runat_11032003_101610_27486/app_query.fasta_1.3576
-DB-GenEmbl -QFMT-fastap -SUEFIX=p2n.rge -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCONE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE=500| -MINLEN=0 -MAXLEN=2000000000
-USER=US09827854_eCGN_1_1_3008_@runat_11032003_101610_27486 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_|SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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Description

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PN JP 1985118
PN 25-JUN-198
PF 29-NOV-198
PF TERANISHI
KIMURA MASAKO,
PI IKEDA YASU
PC C12N15/00,
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CC hypothetic
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Bukaryota; Metazba; C
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Patant: JP 1985118189-A 1 25
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strandedness: Double;
topology: Linear;
hypothetical: No;
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29-NOV-1983 JP 1983224980
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: 1 (bases 1 to 1110)
: 1 (bases 1 to 1110)
: Teranishi,Y., Matsui,Y., Ikeda,Y. and Kimura,M.
: PRODUCTION OF HUMAN APOLIPOPROTEIN E-LIKE PROTEI

Patent: Jp 1986096997-A/1

CC HUMAN (Homo sapiens)
PN JP 1986096997-A/1

PD 15-MAY-1986

PF 16-CCT-1984 JP 1984216987

PI 16-CCT-1984 JP 1984216987

PI TERANISHI YUTAKA, MANSUI YASUSHI, IKEDA YA:
PC C12P21/00,A61K35/74,A61K37/04,C12N15/00,(C:
PC C12P21/00, A61K35/74,A61K37/04,C12N15/00,(C:
PC C12P21/00, A61K35/74,A61K37/04,C12N15/00,(C:
CC hypothetical: No;
CC hypothetical: No;
CC anti-sense: No;
CC *source: tissue_type=Liver;
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16-OCT-1984 JP 1984216987
16-OCT-1984 JP 1984216987
TERANISHI YUTAKA, MATSUI YASUSHI, IKEDA YASUKO, KIMURA WATSUI YASUSHI YAS
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lAlaGluGluThrArgAlaAr                GGCGGAGGAGACGCGGGCACG	YSG1      AGGA	LuGL	laLe           ACT	luPro       AGCCO	laLe	025	000	n="Ho ="ta:	THE I	an-Ba	a; P.C	Paten 38308	lyThrSerAlaAlaF               GCACCAGCGCCGCCC	algl       rgga	lnIl       AGAT	euAsı       GGA
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luLe      AGCT	erGl	alTh []] TCAC	YrLe      ACCT	lnG1      AGCA	laG1       CAGG				нт	E.S.	; V∈		erAs       GCGA	rpal       GGC	laph       CCTT	alal      TGGC
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Patent: JP 2001017028-A 2 23-JJ
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PD 23-JAN-2001
PF 28-APR-2000 JP 200128919
PR SHINOBU FUJITA, HIROKI HAMJ
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                                     SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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        AGCTGGTTCGAGCCCCTGGTGGAAGACATGCAGCGCCAGTGGGCCGGGCTGGTGGAGAAG
                                                                    AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCCAAG
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720 220 660 200 600

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GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140 ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120 ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal CGGCTGGGCGGGCATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG ACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCCAGCAGACCGAGTGGCAGAGC Length:
Matches:
Conservative:
Mismatches:
Indels: 148 420 40 480 360 100 300 80 240 60 180 120

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85279526
                                                                                                                                                                                                 6 (sites)
Rall,S.C. Jr., Newhouse,Y.M., Clarke,H.R., Weisgraber,K.H., McCarthy,B.J., Mahley,R.W. and Bersot,T.P.
Type III hyperlipoproteinemia associated with apolipoprotein phenotype E3/3. Structure and genetics of an apolipoprotein F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84131952
6199196
                                                                                                                                                                                                                                                                                                                                                                                          5 (bases 577 to 624)
Gill, L.L., Peoples, O.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 J. Biol. Chem. 259 (10),
84212473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wallis,S.C., Rogne,S., Cill,L., Markham,A., Edge,M., Woods,D., Williamson,R. and Humphries,S.

The isolation of cDNA clones for human apolipoprotein E and the detection of apoE RNA in hepatic and extra-hepatic tissues
EMBO J. 2 (12), 2369-2373 (1983)
(6) sites; mutations resulting in type III hyperlipoproteinemia. Apo E is a component of normal human very low density lipoprotein. There are six human apo E phenotypes known to result from a single structural gene, three of the common alleles being epsilon-4,
                                                                                                                                                                          variant
J. Clin
                                                                                                                                                                                                                                                                                                                                                          Gill,L.L., Peoples,O.P., Pearston,D.H., Robertson,F.W., Humphries,S.E., Cumming,A.M. and Hardman,N.

Isolation and characterisation of a variant allele of the gene for
                                                                                                                                                                                                                                                                                                                                                                                                                                 6327682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of a new variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human apolipoprotein E mRNA. cDNA cloning and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Taylor,J.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McLean, J.W., Elshourbagy, N.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Breslow, J.L. Synthesis, i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 J. Biol. Chem. 257 (24)
83082756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Breslow, J.L., McPherson, J., Nussbaum, A.L., Williams, H.W., Lofquist-Kahl, F., Karathanasis, S.K. and Zannis, V.I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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K00396
                                                                       epsilon-3 and variant. [5] epsilon-2 allele.
                                                                                                                                                        J. Clin. Invest.
89198059
                                                                                                                                                                                                                                                                                                                                              human apolipoprotein E
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 355 to 1156)
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lis, S.C., Rogne, S.,
                                                                                                                       corrected in
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                                                                                                                                                                                                                                                                                                                              Biophys.
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                                                                                                     ted in [J. Biol. Chem. 258, 11422-11422 (1983)]. 258, 11422-11422 (1983)] correction of [1]. [4]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to 1156)
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(APOE)
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NA, complete o
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/replace="a" 409

citation=[4] /gene="APOE" replace="a" /citation= 'gene="APOE" /replace="a"

note="in variant"

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11422-11422 (1983)],[3]"

/gene="APOE"

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/gene="APOE"

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11422-11422 (1983)],[3],

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Apo E '
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the Apo C-II gene (see separate entry).

The two mutations causing type III hyperlipoproteinemia (apolipoprotein phenotype E3/3) produces substitutions of A (spoilpoprotein E phenotype E3/3) produces substitutions of A Cys at amino acid I12 and Cys for Arg at amino acid position Draft entry and printed copy of sequence for [1] kindly proving the company of the company of the copy of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          [J. Biol. Chem. 258, 11422-11422 (1983)] is too extensively revised relative to [1] to record the revisions in the FEATURES table. The sequence below is that of [J. Biol. Chem. 258, 11422-11422 (1983)]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      see the separate entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Complete source information:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PNAS USA 81,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human liver [1], [J. Biol. Chem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.L. Gill, University of Aberdeen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4] and blood
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                                                                                                                                                                                                                                                                                                                                                                                            /translation="mkvlmaallvTflaGCQAKvEQAVETEPEPELRQQTEWQSGQRW ELALGERWDYLWVQTLSEQVQEELLSSQVTQELRALMDETMKELKAYKSELERGLTP VAEETRARLSKELQAAQARLGADCGREDCGRTUQYRGEVQAMLGQSTEELRVRLASHLE KLRKRLLRDADDLQKRLAVYQAGAREGAERGLSAIRERLGPLVEQGRVRAATVCSLAG QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product="preapolipoprotein
/protein_id="AAB59546.1"
/db_xref="GI:178851"
                                                         /gene="APOE"
                                                                                                                   /note="apolipoprotein
115. .1011
                                                                                                                              115.
                                                                                                                                                                                                                                                                                                                                              ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH"
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1. .1156
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/db_xref="taxon:9606"
/map="19q13.2"
/product="apolipoprotein E
                                                                                                                                                                                                                     /gene="APOE"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="GDB:G00-119-691"
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mature peptide"
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be linked to
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                    NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC
CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Cons
DNA Sequencing by: Genome Sequence Centre,
BC Cancer Agency, Vancouver, BC, Canada
                                                                                                                                                   Submitted (20-FEB-2001) National Institutes of Gene Collection (MGC), Cancer Genomics Office, Institute, 31 Center Drive, Room 11A03, Betheso
                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 4557324
Location/Qualifiers
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/protein_id="AAH03557.1"
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VAEETRARLSKELQAAQARLGADMEDVGELTVQYKGEVQAMMGQSTEELRVERLASHLR
KLRKRLLRDADDLOKELAVYQAGAREGAERGLSAIRERLGPLVEQGRVRAATVGSLAK
KLRKRLLRDADDLOKELAVYQAGAREGAERGLSAIRERLGPLVEGGRVRAATVGSLAG
QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
ARKKSWFEPLVEDMQRCWAGIVEKVQAAVGTSAAPVESDNH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="Eye, retinoblastoma"
/clone_lib="NIH_MGC_16"
/lab_host="DH10B.F"
/note="Vector: pOTB7"
/46._999
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/db_xref="LocusID:348"
/db_xref="Laxon:9606"
/clone="MGC:1571 IMAGE:3355712"
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                                   METHOD FOR PRODUCING APPLIPOPROTEIN
METHOD FOR PRODUCING APPLIPOPROTEIN
PATENTESI OR P 1994315392-A 1 15-NOV-1994;
MITSUBISHI KASEI CORP
OS HOMO SAPIENS (human)
PN JP 1994315392-A/1
PD 15-NOV-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF 21-JUN-1985 JP 1994015433
PF 31-JUN-1985 JP 1994015433
PF 31-JUN-1994 JP 19940154333
PF 31-JUN-1994 JP 19940154333
PF 31-
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MORIMOTO HIRONORI, TERANISHI YUTAKA
C12P21/02,C70K013/00,C12N5/10,C12N15/12,C12N15/18,(C12P21/02,C12R1:91),
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MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/1
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128
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Apo E humanized
BD004277
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JP 2001017028-A/1.
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/db_xref="taxon:9606"
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                                                Scores:
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Sequence 378
AX333278
AX3333278.1
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Patent:
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Eukaryota;
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Cancer gene
                                                                                                                                            Young, P.E., Augustus, M., Horrigan, S., Soppet, D.R.
                                                                                                                Avalon
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                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
370 c 426 g 14
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Unclassified.

1 (bases 1 to 1157)
Attie,A.D., Gretch,D.G., Sturley,S.L. an Production of recombinant proteins in ir patent: US 5472858-A 1 05-DEC-1995;
Location/Qualifiers
1. .1157
ce
                            GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                          GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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6327682
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1 (bases 1 to 1157)
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212473
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                                                                                                                                                                               new variant
                                                                                                                    Location/Qualifiers
1. .1157
  /gene="APOE"
<1. .1157
                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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| . .1157
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/sex="male" |
                                                                                  /map="19q13.2"
                                             tissue_type="liver"
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Best Local Similarity:
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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu 100
                       CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
                                   LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80
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/note="g in pHAE[112,178]; a
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/note="t in pHAE[112,178];
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/note="a in pHAE[112,178];
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/replace="g"
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/product="apolipoprotein
/note="G00-119-691"
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/protein_id="AAB59518.1"
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/db xref="GDB:GOO-119-691"
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/translation="MKVLWAALLVTFLAGCQAKVEQAVETEPEELRQTEWQSGQRW
ELALGREWDYLRWVQTLSEQVQEELLSSQVTQELRALMDETWKELKAYKSELEQLTP
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VAEETRARLSKELQTAQARCAAMEDVGGRLVQYKGFLVQYKGAMATVGSLAG
KLRKRLLRDDDDLQKRLAVYQAGAREGAERGLGFLYQXKGFLYAFTGAAATVGSLAG
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/note="apoE mRNA;
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Apo E humanized mammal
Patent: JP 2001017028-A 3 2
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/3
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128-
PF 28-APR-2000 JP 2000128-
PR SHINOBU FUJITA, HIROKI I
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1156)
Fikni.Y. and Yokoyama, M.
          PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE AOLIKÓT/027, A61K45/00, A61P25/28, A61P43/00,C12N5/10, PC C12N15/09//C07K14/775, PC (C12N5/10,C12R1:91),C12N5/00,C12N15/00,(C12N5/00,C1
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BD004279
BD004279.1 GI:18632240
JP 2001017028-A/3.
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                              SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
                                                                    LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
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Location/Qualiflers
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/db_xref="taxon:9606"
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                     US-09-827-854-15 (1-317) x MFAPOE
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JOURNAL
MEDLINE
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TITLE
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MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
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Macaca fascicularis.
Macaca fascicularis
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
1 (bases 1 to 1178)
                                                                                                                                                                                                                                                                                                                                                                                                                                             Data
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Direct Submission
Submitted (16-DEC-1988) Marotti K.R.,
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X13887
X13887.1 GI:38054
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Location/Qualifiers
                                                                                                                                                                                             83. .136
137. .103
                                                                                                                                                                                                                           /db_xref="SWISS-PROT:P10517"
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                                                                                                                                               /product="apolipoprotein
1154..1159
/note="polyA signal"
373 c 426 g 162
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/db_xref="GI:38055"
                                                                                                                                                                                                                    ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGASTAPVPIDNH"
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AF261279
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                                                                                             dq
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                                                                                complete
 Vertebrata;
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JOURNAL
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AUTHORS
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Nickerson, D.A.
Nickerson, D.A.
Direct Submission
Submitted (27-APR-2000) Department of Molecular Biotechnology,
University of Washington, Box 357730, Seattle, WA 98195, USA
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 5491)

Nickerson,D.A., Taylor,S.L., Fullerton,S.M., Weiss,K.M.,

Clark,A.G., Stengard,J.H., Salomaa,V., Boerwinkle,E. and Sing,C.F.

Sequence diversity and large-scale typing of SNPs in the human apolloportein E gene

Genome Res. 10 (10), 1532-1545 (2000)
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CCTGGGGCCCCTGGTGGAACAGGGCCGCGTGCGGGCCGCCACTGTGGGCTCCCTGGCCGG	4143 CCTGGGGCCCCTGGTGGA	Db
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AGTGTACCAGGCCGGGGCCCGCGAGGCCCCGAGCGCCTCAGCGCCATCCGCGAGCG	4083 AGTGTACCAGGCCGGGGCC	В
aValTyrGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluAr	178 aValTyrGlnAlaGlyAla	Qy
CCTGCGCAAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGC	4023 CCTGCGCAAGCTGCGTAAC	ф
sLeuArgLysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAl	158 sLeuArgLysLeuArgLys	Qу
CGAGGTGCAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCA	3963 CGAGGTGCAGGCCATGCTC	Ър
yGluValGlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHi	138 yGluValGlnAlaMetLe	Qy
GCAGGCCCGGCTGGGCGCGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGG	3903 GCAGGCCCGGCTGGGCGCC	B

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-Q=/cg12_1/USPTO_spool/US09827854/runat_11032003_101610_27486/app_query.fasta_1.3576
-DB=/cenembl -QFWT=fastap -SUFFIX=p2n.rge -MINMATCH+0.1 -LOOPCL=0 -LOOPEXY=0
-UNITS-bits -START=1 -END--1 -MATRIX=blosum62 -TRANS-human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MXX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTPMT=pto -NORM-ext -HEARSIZE=500 -MINLEN=2000000000
-USER=US09827854_eCGN_1_1_13008_@runat_11032003_101610_27486 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPDEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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## SUMMARIES

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PATENT: JP 1985118189-A 1 2:

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PD 25-JUN-1985

PF 29-NOV-1983 JP 1983224:

PI TERANISHI YUTAKA, TAKAI

KIMURA MASAKO,

PI IKEDA YASUKO

PC C12N15/00,CO7H21/04//C:

CC strandedness: Double;

CC topology: Linear;

CC hypothetical: No;

CC antl-sense: No;

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C12N15/00,C07H21/04//C12P21/00;
strandedness: Double;
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25-JUN-1985
29-NOV-1983 JP 1983224980
TERANISHI YUTAKA, TAKAMATSU NOBUHIKO,
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/product='apolipoprotein
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human apolipoprotein
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                                                                       /product='apolipoprotein
                                         /product='apolipoprotein E3 signal peptide' .965
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               LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
                                               SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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/db_xref="taxon:9606"
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1 (bases 1 to 1110)

2 1 (bases 1 to 1110)

Teranishi, Y., Matsui, Y., Ikeda, Y. and Kimura, M.

PRODUCTION OF HUMAN APOLIPOPROTEIN E-LIKE PROTEIN

PAtent: JP 1986096997-A 1 15-MAY-1986;

MITSUBISHI CHEM IND LTD

OS Human (Homo sapiens)

PN JP 1986096997-A/1

PD 15-MAY-1986

PF 16-CCT-1984 JP 1984216987

PF 16-CCT-1984 JP 1984216987

PF C (12PA15/00, A61K35/74, A61K37/04, C12N15/00, (C12P PC (C12N15/00, K11-80))

CC strandedness: Double;

CC topology: Linear;

CC hypothetical: No;

CC *source: tissue_type=Liver;

FH Key

FT sig_peptide J5..68

FT sig_peptide /product='human apolipoprotei
GluGlnalaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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                                                GAGCAAGCGGTGGAGACAGAGCCGGAGCCGAGCCAGCAGACCGAGTGGCAGAGC
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15-MAY-1986
16-OCT-1984 JP 1984216987
TERANISHI YUTAKA, MATSUI YASUSHI, IKEDA YASUKO, KIMURA MASAKO TERANISHI YUTAKA, MATSUI YASUSHI, IKEDA YASUKO, KIMURA MASAKO C12P21/00,A61K35/74,A61K37/04,C12N15/00,(C12P21/00,C12R1:19),
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/db_xref="taxon:32644"
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                                                   human.
Homo sapien
Eukaryota;
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Sequence 63
AX302545
AX302545.1
                    Tumor markers in ovarian cancer Patent: WO 0175177-A 63 11-OCT-2001; THE SECRETARY OF THE DEPARTMENT OF H
                                    Morin, P
                                               Mammalia;
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                              markers in ovarian car
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Eutheria;
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                                              Chordata;
Primates;
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WO0175177.
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              Patent: JP 2001017028-A 2 23-JAN-2001;
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
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PD 23-JAN-2001
PF 28-APR-2000 JP 2000128919
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PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO
                                                                         LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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A01K67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC
C12N15/09//C07K14/775,
PC (C12N5/10, C12R1:91), C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
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Fujita,S., Hamanaka,H., Fukui.
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                                                                                 Wallis,S.C., Rogne,S., Gill,L., Markham,A., Edge,M., W
Millianson,R. and Humphries,S.
The isolation of cDNA clones for human apolipoprotein
detection of apoE RNA in hepatic and extra-hepatic tis
EMBO J. 2 (12), 2369-2373 (1983)
Breslow, J.L.
Synthesis, intracellular
apolipoprotein E
                                                                                                                                                                                                              Breslow,J.L., McPherson,J., Nussbaum,A.L., Williams,H.W., Lofquist-Kahl,F., Karathanasis,S.K. and Zannis,V.I. Identification and DNA sequence of a human apolipoprotein
                                                                                                                                                                                                                                                 Homo sapiens
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Zannis, V.I., McPherson, J.,
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[J. Biol. Chem. 258, 11422-11422 (1983)] is too extensively revised relative to [1] to record the revisions in the FEATURES table. The sequence below is that of [J. Biol. Chem. 258, 11422-11422 (1983)]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biochem.
85279526
                                                                                                                                                                                                                                                                                                                                         Apo E is located on chromosome 19 --Jackson, Bruns and Breslow , PNAS USA 81, 2945-2949 (1984) -- and is believed to be linked to the Apo C-II gene (see separate entry).

The two mutations causing type III hyperlipoproteinemia (apolipoprotein E phenotype E3/3) produces substitutions of Arg for Cys at amino acid 112 and Cys for Arg at amino acid gosition 142.

Draft entry and printed copy of sequence for [1] kindly provided by L.L. Gill, University of Aberdeen.

Complete source information:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             J. Biol.
84185684
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Rall, S.C. Jr., Newhouse, Y.M., Clarke, H.R., Weisgraber, K.H., McCarthy, B.J., Mahley, R.W. and Bersot, T.P.
Type III hyperlipoproteinemia associated with apolipoprotein phenotype E3/3. Structure and genetics of an apolipoprotein E
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epsilon-3 and variant.
[5] epsilon-2 allele.
[6] sites; mutations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Taylor, J.M.
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                                                                      /gene="APOE"
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                                                                                                                                         /tissue_type="liver and blood"
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QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
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SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
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                                                                                                                                                                                                          AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Contact: MGC help desk
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info@bcgsc.bc.ca
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/db_xref="LocusID:348"
/db_xref="taxon:9606"
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/tissue_type="Type, retinoblastoma"
/clone_lib="NHL_MGC_16"
/lab_host="DH10B-R"
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                                    LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
                                                                          ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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Patent: JP 1994315392-A 1 15-NOV-1994;
MITSUBISHI KASEI CORP
OS HOMO Sapiens (human)
PN JP 1994315392-A/1
PD 15-NOV-1994
PP 11-JUN-1985 JP 1994015433
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PF (12P21/02,CO7K13/00,C12N5/10,C12N15/PP C (12R1:91),
PC (12R1:91),
PC (12R1:91),
PC (12R1:91);
CC strandedness: Double;
CC topology: Linear;
FH Key Location/Qualifiers
FH Key Location/Qualifiers
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FT sig_peptide 15..68
FT mat_peptide 69..968
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/organism='Homo s
g_peptide 15..68
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         Similarity:
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Sequence 378
AX333278
AX3333278.1
                                                                                                       Patent:
Avalon I
                                                                                                                            Young, P.E., Augustus, M., Carte
Horrigan, S., Soppet, D.R. and V
Cancer gene determination and
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                      WO 0194629-A 3787 13-DEC-2001;
Pharmaceuticals (US)
Cocation/Qualifiers
1. .1157
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/db_xref="taxon:9606"
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/note="EMBL/GenBank Accession
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Sequence
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1 (bases 1 to 1157)
Attie,A.D., Gretch,D.G., Sturley,S.L.
Production of recombinant proteins in
Patent: US 5472858-A 1 05-DEC-1995;
Location,Qualifiers
1. 1157
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                                                                                                                                                                                                                                     Taylor,J.M.
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 /gene="APOE"
<1. .1157
                           /tissue_type="liver"
/dev_stage="57-year old"
1. .1157
                                                                 /map="19q13.2"
/clone="pHAE[112,178,813]."
/sex="male"
                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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/product="apolipoprotein
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/protein_id="AAB59518.1"
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/codon_start=1
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                                         Fujita, S., Hamanaka, H., Fukui, Y. and Yokoyama, M.
Apo E humanized mammal
Patent: JP 2001017028-A 3 23-JAN-2001;
MITSUBISHI CHEMICAL CORP
OS Homo saplens (human)
PN JP 2001017028-A/3
PD 23-JAN-2001
PF 28-APR-2000 JP 200128919
PR SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINI
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; E

Mammalla; Eutheria; Primates; Catarrhini; Hominidae;

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                                                                                                                                                                                                                                                                                                            ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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RESULT 14
MFAPOE
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MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20
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Nucleotide sequence of the cynomolgus monkey apolipoprotein E cDNA
Nucleic Acids Res. 17 (4), 1778 (1989)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Direct Submission
Submitted (16-DEC-1988) Marotti K.R., The Upjohn Co,
301 Henrietta Street, Kalamazoo, MI 49008
2 (bases 1 to 1178)
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1 (bases 1 to 1178)
Marotti, K.R.
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Macaca fascicularis.
Macaca fascicularis
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Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
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1154..1159
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373 c 426 g 162
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83. .1036
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AF261279 5491 bp DNA linear PRI 27-OCT- HOmo sapiens apolipoprotein-E gene, complete cds. AF261279 AF261279.1 GI:11034800 Homo sapiens Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom	ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 317                 	SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300 	LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 280 	SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys 260	LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240 	ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220 	GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200 	LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr 180 	GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160 	ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140 	ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120 :::	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluGlnLeu 100 	LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80 	GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 60 	GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40	
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1. .5491
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Nickerson, D.A.
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Genome Res. 10 (10), 1532-1545 (2000)
20499366
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 5491)
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73
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118 aGlnAlaArgLeuGlyAlaAspMetGlu 	98 uGlnLeuThrProValAlaGluGluThrA 	79 -ArgalaLeuMetaspGluThrMetLys 	3723 CCCCATCCCAGCCCTTCTCCCCGCCTCC	78	3663 GCCTCTGCCCCGTTCCTTCTCTCTCTCT	78	3603 TCGTCTCTGCCTCTGCCTCTGCATCTG	78	3543 CCTCCCAAAGTGCTGGGATTAGAGGCAT	78	3483 CTCACTGTGTTGCCCAGGCTGGTCTTGA	78	3423 CGTCCTGGCTCTGTCTCTGTCCTTCCCT	78	3363 TGGAATTCTCTCTCTCAGCTTTGTCTCT	78	3303 CCTGGGTCTCTGCTGGTTCTAGCTTCCT	78	3243 GGCGGCTATACCTCCCCAGGTCCAGGTT	78	3183 TCCCAGGTCACCCAGGAACTGAGGTGAG	erg	52 TrpAspTyrLeuArgTrpValGlnThrI 	3 CTGCGCCAGCAGACCGAGTGGCAGAG	32 LeuArgGlnGlnThrGluTrpGlnSe	12 PheLeuAlaGlyCysGlnAlaLysValo		ignment Scores:  ed. No.:  4.32e-71  2.72e-71  2.72e-71  2.72e-71  2.72e-71  2.72e-71  2.72e-71  2.72e-71  2.72e-71  2.72e-71  3.72e-71  3.72e-71  3.72e-71  3.72e-71  4.32e-71  3.72e-71  4.32e-71  60.608  60.608  60.608  87.388  67.388
.uAspValCysGlyArgLeuValGlnTyrArgGl 1 	rArgAlaArgLeuSerLysGluLeuGlnAlaAl 1 	SGluLeuLysAlaTyrLysSerGluLeuGluGl 9 	CACTGTGCGACACCCTCCCGCCCTCTCGGCCG 3	7	FTGGGTCTCTGGCTCATCCCCATCTCGCCCG 3	7	3CTCTCTGCATCTGTCTCTGTCTCCTTCTCG 3	7	REAGCCACCTTGCCCGGCCTCCTAGCTCCTTCT 3	7	AACTTCTGGGCTCAAGCGATCCTCCCGCCTCGG 3	7	PAGCTCTTTTATATAGAGACAGAGAGATGGGGT 3	7	TCTCTCTTCCCTTCTGACTCAGTCTCTCACACT 3	7	TCTTCCCATTTCTGACTCCTGGCTTTAGCTCTC 3	7	TTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGGG 3	7	STGTCCCCATCCTGGCCCTTGACCCTCCTGGTG 3	7	nThrLeuSerGluGlnValGlnGluGluLeuLeuSer 7	CTGGGAACTGGCACTGGGTCGC	rGlyGlnArgTrpGluLeuAlaLeuGlyArgPhe 5	GluGlnAlaValGluThrGluProGluProGlu 3 	1-54,91)	Length: 5491 Matches: 303 Conservative: 0 Mismatches: 3 Indels: 194 Gaps: 1
L38	118 3902	98 3842	3782	78	3722	78	3662	78	3602	78	3542	78	3482	78	3422	78	3362	78	3302	78	3242	78	71 3182	3122	51	31 3062		

278 gLeuLysSerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuVa
258 gAlaLysLeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaAr 
238 umetGlySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValAr 
218 yGlnProLeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGl 
198 gLeuGlyProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGl 
178 aValTyrGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluAr 
158 SLeuArgLySLeuArgLySArgLeuLeuArgAspAlaAspAspLeuGlnLySCySLeuAl 
138 yGluValGlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHi 
3903 GCAGGCCCGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGG

Search completed: March 14, 2003, 17:32:29 Job time: 2314.23 secs

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MODEL-frame+_p2n.model -DEV=xlp

-Q-cgn2_1/USPTO_spool/US09827854/runat_11032003_101609_27476/app_query.fasta_1.3576
-D=N_Geneseq_101002 -QFMT=fastap -SUFFIX=p2n.rng -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -ENb=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST-45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=sct -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
-USER=US09827854_eCGN_1_1_1201_@runat_11032003_101609_27476 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEC_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPDEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1993.DAT:*

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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1000.DAT:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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## SUMMARIES

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AAQ11980 AAC02139 AAH98479 AAV39612	ABL62679 ABL67340 AAS03049	AAS22673 AAF18114 AAV89595 ABK34238 ABN95994	AAQ69101 AAT18070 AAT69792 AAT18068 AAQ69099	AAZ09524 AAZ09526 AAD26108 AAD32081 AAD32077 AAD32075 AAD32075 AAV29159	AAF84314 AAD22050 AAD22035 AAN60409 AAD26035 AAN60409 AAB83113 AAF84315 AAD22052 AAD22052 AAD22051 AAT06957 ABN95746 ABK64514 ABL65450 AAF84316 AAD22047 AAR822437 AAX75756 ABL31915	ID
Human apolipoprote Human secreted pro Human EST-derived E2 allele DNA enco	Colon adenocarcino Thyroid cancer rel Human diagnostic a	Human CDNA encodin Lung cancer associ EST clone CP147. Human CDNA for nov Gene #2407 used to	ApoE4Lx2 protease Human ApoE4Lx2 cDN Partial human apol Human ApoE4 CDNA. ApoE4L protease cD	Human Apo E genomi Human Apo E genomi Human apolipoprote Bovine ApoE gene. Human alpha-1-anti Human albumin prom Nucleotide sequenc	APOEZ cc apolipop bearign g cancer re can	Description

## ALIGNMENTS

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RESULT 1
AAF84314
ID AAF84314;
XX

AC AAF84314;
XC AAF84314;
XX

DT 21-JUN-2001 (first entry)
XX

DE Human ApoE2 coding sequence.
XX

KW Human; ApoE2; Alzheimer's disease; arteriosclerosis; ss.
XX

CS Homo sapiens.
XX

CS Homo sapiens.

EH Key Location/Qualifiers
FT CDS 61..1014
FT /*tag= a
/*tag= a
/*product= "Human ApoE2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an ApoE humanised mammalian cell. The present sequence is the coding sequence for human ApoE2, which was use in the method of the present invention. The ApoE humanised mammalian can be used for screening for agents useful for treating or preventing Alzheimer's disease and arteriosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New apoE humanized mammalian cell useful for screening fo useful for treating or preventing Alzheimer's disease and {\tt arteriosclerosis} -
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                                       CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCCACCTGCGC
                                                                               CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
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hypertriglyceridaemia; low density lipoprotein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cholesterol, delaying the onset of or treating mammal, without inducing hypertriglyceridemia
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          LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
                                            ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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                                CAGGCCGGGGCCCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
                                                                                                                          AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGTGCCTGGCAGTGTAC
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                                                           New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia -
                                                                                                                                             06-APR-2000; 2000US-0544386
04-OCT-2000; 2000US-0679088
05-APR-2001; 2001US-0827854
                                 The present sequence is a human apolipoprotein E (apoE)
                                               Claim 14;
                                                                                       WPI; 2002-010885/01.
P-PSDB; AAE13296.
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allele, apoEl DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (I

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                                                                            GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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Ct tumours in an individual via the detection and measurement of the cexpression of ovarian tumour marker genes (ABA83081-ABA83122, ABA83180, CC ABA83182 and ABA83184) or segments thereof (ABA83081-ABA83122, ABA83199, CC ABA83182 and ABA83184) or segments thereof (ABA83181 and ABA83184) or segments thereof (ABA83181 and ABA83189). The methods of the invention are useful for CC detecting an ovarian tumour in a patient, for identifying an individual CC at increased risk for developing ovarian cancer, in prognostic tests for CC monitoring a patient in remission from ovarian cancer and in tests for CC monitoring a patient in a patient being treated for ovarian cancer. The methods can additionally be used to identify a particular tumour as Serous cystadenoma, borderline serous tumour, serous cystadenocarcinoma, concinous cystadenoma, borderline serous tumour, mucinous cystadenoma, borderline serous tumour, mucinous cystadenocarcinoma, cystadenofibroma, adenofibroma and Brenner CC tumour. The ovarian tumour marker genes of the invention were identified cusing SAGE (serial analysis of gene expression) and were found to be coverexpressed in a broad variety of ovarian epithelial tumour cells cells crelative to normal ovarian epithelial cells. The marker genes are CC inplicated in immune response pathways, in the regulation of cell coalised or secreted. In addition to their use as diagnostic and CC proteins may be used as therapeutic targets for the treatment and coal coal coal coal coal cancer. Sequences ABA83081-ABA83182, ABA83180, CC ABA83182 and ABA83184 represent the ovarian tumour marker genes of the invention.
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                                                                                                                                                                                                                                                                       present sequence is the coding sequence for in the method of the present invention. The can be used for screening for agents useful Alzheimer's disease and arteriosclerosis.
                                                                                                                                                                                                                                                                                                                                      New apoE humanized mammalian cell useful for screening useful for treating or preventing Alzheimer's disease arteriosclerosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDS
                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                      Disclosure;
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                                   LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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DB; AAB80997.
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                                                                                                                                                                                                                            sig_peptide
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hypertriglyceridaemia; low density lipoprotein; LDL; ds.
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(UYBO-)
               WPI; 2002-010885/01
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                                             KOS P
                                                                  ; 2000US-0544386.
; 2000US-0679088.
; 2001US-0827854.
                             Kypreos
                                                    PHARM INC.
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density lipoprotein; LDL; ds.
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PT New apolipoprotein E polypeptide and nucleic acid, useful for lowering PT cholesterol, delaying the onset of or treating atherosclerosis in PT mammal, without inducing hypertriglyceridemia - XX Claim 14; Page 83; 91pp; English.
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The present sequence is a human apolipoprotein E (apoE) allele, apoE2\*\* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LDL) receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or having a defect in remnant removal.

Sequence 1156 BP; 207 A; 369 C; 432 G; 148 T; 0 other;

Alignment Scores:

Pred. No.:

9.56e-115

Core:

1573.00

Matches:

1573.00

Matches:

1573.00

Matches:

1573.00

Matches:

1

Conservative:

1

Best Local Similarity:

99.37%

Buest Local Similarity:

98.99%

Conservative:

1

Mismatches:

0

Caps:

Caps

US-09-827-854-16 (1-317) x AAD22052 (1-1156)

240	LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly	221	Qy
720		661	Db
220	ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro	201	Qy
660	CAGGCCGGGGCCCGCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG	601	Db
200		181	Qy
600		541	Db
180	LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr	161	Qy
540		481	Db
160	GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg	141	Qy
480	CGGCTGGGCGCGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG	421	ДĎ
140		121	Qу
420		361	Db
120	ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla	101	Qy
360		301	Db
100	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu	81	Qy
300		241	DЬ
80	LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla	61	Qy
240	GCCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA	181	рь
60		41	Qy
180	GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC	121	рь
40		21	Qy
120	ATGAAGGTTCTGTGGGCTGCGTTGCTCGCTCACATTCCTGGCAGGATGCCAGGCCAAGGTG	61	Db
20		<u>_</u>	Qy

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04-OCT-2000;
05-APR-2001;
The present sequence is a human apolipoprotein E (apoE) allele, apoE2* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous.
                                                                                              Claim
                                                                                                                      New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia
                                                                                                                                                                                                                                                            (KOSP-)
(UYBO-)
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hypertriglyceridaemia;
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»B; AAE13297.
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normally functioning apoE gene or low density lipoprotein receptor or is at risk for developing atherosclerosis due accumulation of lipoprotein remnants in the bloodstream or a defect in remnant removal.
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                                                                                                                                                                                                                                                                                                                                                                                                     Prodn. of recombinant apo:lipoprotein E in insects - by infect Manduca sexta larvae with recombinant Autographica californica nuclear polyhedrosis baculovirus vector.
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                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method for diagnosing and detecting the CC progression of liver cancer, hepatocellular carcinoma or metastatic liver ctumour in a patient, and differentiating metastatic liver cancer from CC hepatocellular carcinoma in a patient, involving detecting the level of expression of two or more genes represented in ABN93503-ABN97455 in a CC tissue sample. The method of the invention has hepatocropic, and CC typostatic activity. The method is useful for diagnosing and detecting CC liver carcinoma in a patient. The method is useful for identifying CC expression profiles which serve as useful diagnostic markers as well as CC markers that can be used to monitor disease states, disease progression, CC drug toxicity, drug efficacy and drug metabolism. CC drug toxicity, drug efficacy and drug metabolism. CC specification, but was obtained in electronic format directly from WIPO CC at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
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tumour; cytostatic; expression profile; disease state;
ion; drug toxicity; drug efficacy; drug metabolism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of diagnosing (I) the onset or progression of benign prostatic hyperplasia (BPH), or screening (II) for or identifying an agent that modulates the onset or progression of BPH.

CC The method is based on changes in gene expression in BPH tissue isolated from patients exhibiting different clinical states of prostate compared to normal prostate tissue. (I) comprises CC detecting the expression levels of one or more genes in prostate cells composed that are differentially regulated compared to normal composite that are differentially regulated compared to normal composite cells. (II) comprises preparing a first gene expression profile constate cells. (II) comprises preparing a first gene expression profile cells, and comparing the cell population, exposing the cells to the cells, and comparing the first and second gene expression profiles. (I) is useful for diagnosing the onset or progression profiles. (I) is useful for diagnosing the onset or progression of BPH. (II) is confiled for identifying an agent that modulates the onset or progression confidentifying confidentifying an agent that modulates the onset or progression confidency is the expression level in a tissue or cells, by comparing the expression confidency in the diabase, and displaying the composite of the expression levels of at least one gene in the tissue or cells sample composited to the composite of the composite of the database, and displaying the composite of the composite of the composite of the database.
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                                                                ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
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RESULT 14
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ABL65450

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The present invention describes a method (MI) for screening for an anti-neoplastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change in expression of at least one gene (I) of a signature gene set, where (I) comprises a sequence (S) selected from 8447 sequences (given in ABL61664 to ABL70110), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has cytostatic activity and can be used in gene therapy. MI can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a cresult of MI, and the data is sufficient to convey the chemical structure and/or properties of the agent. MI can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, cosophageal, ovarian, kidney, prostate or pancreatic cancer, infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.
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Best Local Query Match DB:

Match:

Similarity

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Length:
Matches:
Conservative:
Mismatches:
Indels:

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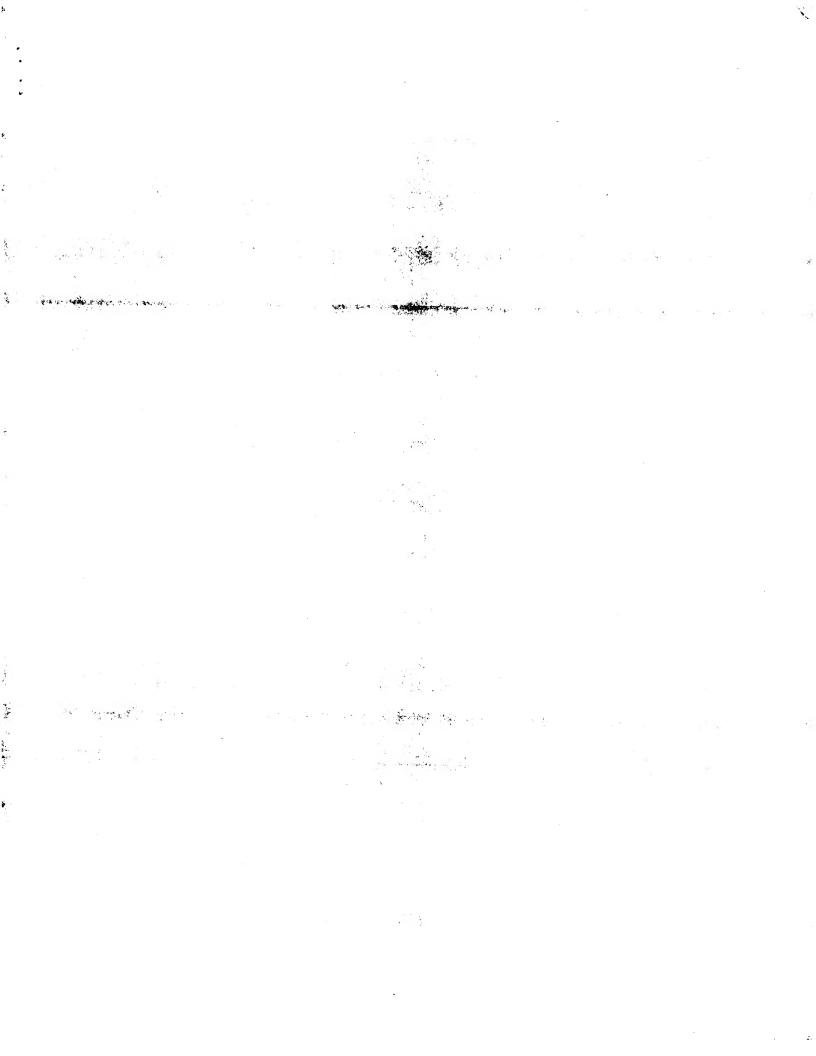
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OY 121 ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140	Oy 101 ThrProValA1aGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120	Oy 81 LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluGlnLeu 100 	Oy 61 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80	Oy 41 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 60	Qy 21 GluGlnAlaValGluThrGluPrOGluPrOGluLeuArgGlnGlnThrGluTrpGlnSer 40	Qy 1 MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20 	US-09-827-854-16 (1-317) x AAN50450 (1-1110)	Alignment Scores: 2.7e-114 Length: 1110 Score: 1567.00 Matches: 314 Percent Similarity: 99.05% Conservative: 0 Best Local Similarity: 99.05% Mismatches: 3 Query Match: 98.62% Indels: 0 DB: 66	SQ Sequence 1110 BP; 198 A; 355 C; 414 G; 143 T; 0 other;	CC The sequence may be used to produce the apolipoprotein E, useful in the CC treatment of hyperlipaemia.	PS Claim 3; Page 484; 8pp; Japanese. XX	PT DNA fragment - contg. DNA which codes human apolipoprotein E for PT treatment of hyperlipaemia.	DR WPI; 1985-188003/31. DR P-PSDB; AAP51204.	PA (MITU) MITSUBISHI CHEM IND KK.	PR 29-NOV-1983; 83JP-0224980.	PF 29-NOV-1983; 83JP-0224980.	PD 25-JUN-1985.	AA PN JP60118189-A. XX	FT mat_peptide 69.965 FT mat_peptide 69.965 FT /*tag= b	Key Location CDS 15968	OS Homo sapiens.	XX XX XX	XX DE Sequence encoding human apolioprotein E.	
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261 LeuGluGluGlnAlaGlnGlnTleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 280	2у 26 Db 79
241 SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys 260	ОУ 24
	ОБ 73
221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240	Ду 2;
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201 ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220	Ωy 2(
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181 GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200	Ωy 18
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141 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160 	Qy 1.

Search completed: March 14, 2003, 12:43:10 Job time : 183.728 secs



Sequence Sequence Sequence

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-DB=Issued_patents_Na -QFMT=fastap -SUFFIX-p2n.rni -MINMARCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX-blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
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-NO_XLPXY -NO_MMAP -LARGEQUERY -NEGL_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
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S-08-726-306A-28

S-08-726-24

S-09-261-256-24

S-09-261-256-24

S-09-274-245-24
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US-07-709-949-1
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                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/07709949 Patent No. 5472858
                                                                                                     ADDRESSEE: Quarles & Brady
STREET: P.O. Box 2113
CITY: Madison
STATE: WI
COUNTRY: USA
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, v
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/07/709,949
FILING DAYE: 19916604
CURASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
     INFORMATION FOR SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Attie, Alan D
APPLICANT: Gretch, Daniel G
APPLICANT: Sturley, Stephen L
APPLICANT: Beckage, Nancy E
TITLE OF INVENTION: Production of
TITLE OF INVENTION: Insect Larvae
                                         REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
                                                                ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                            TELEPHONE:
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US-08-952-796-1
US-08-938-105-2
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US-08-968-751-3
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US-08-961-527-126
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PCT-US93-06160-3
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Sequence

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SEQUENCE CHARACTERISTICS: LENGTH: 1157 base pair

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Best Local Similarity:
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FEATURE:
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LOCATION: 62..119
PUBLICATION INFORMATION:
AUTHORS: McLean, J W
JOURNAL: Journal of Biological Chemistry
VOLUME: 259
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ORIGINAL SOURCE:
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DATE: 1984
RELEVANT RESIDUES
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GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
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                                                                                                                                                                                                                                                                                                                                                                  ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                               TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                 APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1097
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60
FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Piedrahita, Jo
APPLICANT: Bazer, Fuller
TITLE OF INVENTION: Compo
TITLE OF INVENTION: Gener
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                LENGTH:
TYPE: n
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STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK:177
                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/949;155 FILING DATE: Concurrently Herewith
                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: ARNOLD, WHITE AND DURKEE STREET: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
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Generation of Transgenic
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Query Match:
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             ArgLeuLysSerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeu
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                                                             ArgAlaLysLeuGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAla
                                                                                                GAGATGGGCAGCCGGACCGGCCTGGATGAGATGCGTGAGCAGCTGGAGGAGGTG
                                                                                                              GluMetGlySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluVal
                                                                                                                                                                                                             ArgLeuGlyProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAla
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                                                                                                                                                 GGCCAGCCGCTGCGCGAGCGCGCGAAGCCTGGGGCCAGAAGCTGCGCGGACGGCTGGAG
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                                                                                                                                                                                         Query Match:
                                                                                                                                                                                                           Best Local Similarity:
                                                                                                                                                                                                                            Percent Similarity:
                                                                                                                                                                                                                                                             Pred. No.:
                                                                                                                                                                                                                                                                                 Alignment Scores:
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TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
 111
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FILING DATE: «Unknown)
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTARTION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK:177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   939 GTGGAGAGGATGCAGTCGGCCGTGAGCATAAGCTCCTCCACCTCTGCGCCCCAGTGATAAT 998
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                              21 GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu 37
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                     MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20
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ADDRESSEE: ARNOLD,
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                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
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STATE: TX
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                                                                                                                               Sequence 51, Application US/08949155 Patent No. 6271436 GENERAL INFORMATION:
                                                               NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                     TITLE OF INVENTION: Compositions and Methods TITLE OF INVENTION: Generation of Transgenic
                                                                                                          APPLICANT: Piedrahita, APPLICANT: Bazer, Full
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COUNTRY: US
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                                            P.O. Box 4433
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TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TATELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US 60/027,338
FILING DATE: 11-OCT-1996
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APPLICATION NUMBER:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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US-09-819-964-51
; Sequence 51, Application US/09819964
; Patent No. 6369294
; GENERAL INFORMATION:
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               CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WH:
STREET: P.O. Box 4433
                                                                                           APPLICANT: Piedrahita, Jorge
Bazer, Fuller W
                                                      NUMBER OF SEQUENCES: 51
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CITY: Houston
                                                                OF INVENTION: Compositions and Methods Generation of Transgenic
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DB:
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Best Local Similarity:
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TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TA
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 60/046,094 FILING DATE: 09-MAY-1997 ATTORNEY/AGENT INFORMATION:
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                                       CCCTCCTGGCGAACCGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
                                                                                                                                                                                                 TGGCAGGCCAGCCCTGGGAGCAGGCCCTGGGCCGCTTCTGGGATTACCTGCGCTGG
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                                                                                                                                                                                                                                                                                                                                                           ATGGCTGTGAACTGGCTCGGT----
                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: <Unknown>
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COUNTRY: US
ZIP: 77210-4433
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RESULT 6
US-08-726-306A-28
                                                                                  Sequence 28, Application US/08726306A Patent No. 5958684 GENERAL INFORMATION:
              APPLICANT: van Leeuwen, Frederik Willem APPLICANT: Burbach, Johannes Peter Henri APPLICANT: Grosveld, Franklin G. TITLE OF INVENTION: DIAGNOSIS METHOD AND NUMBER OF SEQUENCES: 189
    NUMBER OF SEQUI
                                                                                                                                                                    GCGCCCAGTGATAAT
                                                                                                                                                                                             ValProSerAspAsn 316
                                                                                                                                                                                                                                         GlnTrpAlaGlyLeuValGluLysValGlnAlaAlaValGly---ThrSerAlaAlaPro
                                                                                                                                                                                                                                                                             LeuGlnLysCysLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeu
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                                                                                                                                                                                                                                                                                                                                                                                          US-09-827-854-16 (1-317) x US-08-726-306A-28 (1-660)
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
TELEPAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 660 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/009,832
FILING DATE: 01-Jan-1996
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M
REGISTRATION NUMBER: 34,380
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 95/20080.4
FILING DATE: 02-0ct-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                                                                                                                                                 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                                     ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120
                                                                                                                                                                  CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
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                                                        ACCCCGGTAGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC
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                                                                                                                           Percent Similarity:
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Sequence 6, MPE-
NO. 5525493
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                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 16787/168/DFBC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEPAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/849,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                    MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 330 base pairs
                                                                                                                                                                                                                                         FEATURE:
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APPLICANT: UHLEN, Mathias
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CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                               LOCATION:
                                                                                                                                                                                                                          NAME/KEY:
                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
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AAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTGACCCCGGTGGCGGAGGAG
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1800 Diagonal Road,
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US-09-827-854-16 (1-317) x US-08-617-256-24 (1-252)
                               Query Match:
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                                            Best Local
                                                      Percent Similarity:
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                                                                                   Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. 6043031
GENERAL INFORMATION:
                                                                                                                                                                                     TELEFAX: (617)227-594: INFORMATION FOR SEQ ID NO:
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: SQ
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: March 18,
                                                                                                                     TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION: NAME: Arnold, Beth A.
                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE: March 17.
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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US-09-287-141-24
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                  Pred.
                               Alignment Scores:
                                                               US-09-287-141-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24, Application US/09287141 Patent No. 6197498
                                                                                                                                                                     TELEPHONE: 619-587-5360
TELEPAX: 619-587-5360
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
                                                                                                                                         TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE: 252 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002D
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: % L. STREET: % L. CITY: La Jolla "Mare: CA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor
                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: K ster, Hubert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                  No.:
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                                                                                             STRANDEDNESS:
TOPOLOGY: lir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62
                                                                                                                                TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette
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                                                                                                                           nucleic acid
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                                                                                                                                              252 base pairs
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Length:
Matches:
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Best Local Similarity:
Query Match:
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                                                                              TELEFAX: (617)227-594 INFORMATION FOR SEQ ID NO:
                                                                                                                                                         REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002G
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
                                             SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                                                                                                     FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
                                                                                                                           TELECOMMUNICATION INFORMATION:
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TELEFAX: 619-587-5360
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM CON
OPERATING SYSTEM:
                STRANDEDNESS:
                                                                                                                                                                                                             NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,77
                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/6
FILING DATE: 18-MAR-1996
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 92037-9103
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                                                                                                            TELEPHONE:
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E: CA
                            nucleic acid
                                                                                             : (617)227-7400
(617)227-5941
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Indels:
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Conservative: Mismatches: Indels:

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APPLICANT: K ster, Hubert
TITLE OF INVENTION: DNA D
                         TELEFAX: 858-587-5360
INFORMATION FOR SEQ ID NO:
                                                        NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002J
TELECOMUNICATION INFORMATION:
TELEPHONE: 858-450-8400
                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/6:
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/44
                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                SEQUENCE CHARACTERISTICS:
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                                                                                                                                      FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 42
CITY: La CATATE: CACOUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    149
                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Release #1.0, Version
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7th Floor
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; STRANDEDNESS: sing
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-09-504-245-24
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                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
TITLE OF INVENTION: DNA Diagnostics Based on
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                        ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                               OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                         APPLICATION NUMBER: 08/4
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
REGISTRATION NUMBER: 33,
                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
TELECOMMUNICATION INFORMATION:
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        REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
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                                          Seidman, Stephanie L
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Mass Spectrometry

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TELEPHONE: 619-450-8400 TELEFAX: 619-587-5360

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US-09-287-679-24
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Patent No. 6258538
GENERAL INFORMATION:
APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
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INFORMATION FOR SEQ ID NO:
                                                                                                                                             ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
                                                                                     COMPUTER: IBM COMPATID OPERATING SYSTEM: DOS SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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CITY: La
STATE: CA
COUNTRY:
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TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                             NUMBER OF SEVERITY OF STATE OF THE STATE OF 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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LENGTH: 252 base pairs
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TELECOMMUNICATION INFORMATION:
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NAME: Seidman, Stephanie L
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                                                       COMPUTER: IBM Com
OPERATING SYSTEM:
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COUNTRY: USA
ZIP: 92037-9103
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(617)227-5941
                               PatentIn Release #1.0, Version #1.25
                                                                                              IBM Compatible
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Best Local Similarity:
Query Match:
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                                                                                                                Sequence 24, Application US/09287681
Patent No. 6277573
GENERAL INFORMATION:
APPLICANT: K ster, Hubert
TITLE OF INVENTION: DNA Diagnostics Based on
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSORE: USALIE FRANCE FARMER UNITED TO MAINING FOR
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TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
C11.
STATE: C.
COUNTRY: USA
COUNTRY: 92037-9103
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LENGTH: 252 base pairs
TYPE: nucleic acid
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TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
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APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.:
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                                                                      ADDRESSEE: Heller burners, STREET: 4250 Executive Square,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-20021
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                                                                                                                                                                                                                                                                                                        GluArgGlyLeu 192
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                                                                                        Heller Ehrman White & McAuliffe
250 Executive Square, 7th Floor
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Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                             ; MOLECULE TYPE: cDNA US-09-287-681-24
                                                                                                                                                                                                                                                                                                                                                       Score:
                                                                                                                                                                                                                                                                                                                                                                       Pred.
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TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                    No.:
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                                                                                                       241
                         189 GluArgGlyLeu 192
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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REFERENCE/DOCKET NUMBER: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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 GAGCGCGGCCTC
                                                    GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGCCCGCGAGGGCCGCC
                                                                  AspAlaAspAspLeuGlnLysCysLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla
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Search completed: March 14, 2003, 20:18:40 Job time : 39.7343 secs

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Command line parameters:

-MODEL-frame+_p2n.model -DEV-xlp
-Q-cgn2_1/USPT0_spool/US09827854/runat_11032003_101612_27578/app_query.fasta_1.3576
-DB-Published_App_lsoations_NA -QFMT-fastap -SUFFIX-p2n.rnpb -MINMATCH=0.1
-LOOPCL-0 -LOOPEXT=0 -UNITS-bits -START=1 -END--1 -MATRIX-blosum62
-TRANS-human40.cdi -LIST-45 -DOCALIGN=200 -THR.SCORE-pct -THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE-LCCAL -OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN=0
-MAXLEN=200000000 -USER-US09827854|@CGN_1_1_298_@runat_11032003_101612_27578
-NCPU=6 -ICPU=3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOD=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELEXT=7
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Perfect score:
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Ygapop 10.0 , Ygapext
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                                                                                                                                                                                                                                                                                                                                              /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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Sequence 9, Appli
Sequence 10, Appl
Sequence 129, App
Sequence 17, Appl
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equence 5, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,	Sequence 8, Appli Sequence 454, App Sequence 12, Appl Sequence 11, Appl Sequence 760, App Sequence 2244, Ap Sequence 7, Appl1

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## ALIGNMENTS

RESULT 1 US-09-827-854-9

GENERAL INFORMATION:

Sequence 9, Application US/09827854 Patent No. US20020123093A1

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Percent Similarity:
                                            Pred. No.:
                                                                    Alignment Scores:
                                                                                                                   ; ORGANISM: Homo sapiens US-09-827-854-9
                                                                                                                                                                                                                 SEQ ID NO 9
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without inducing hypertriglyceridemia
FILE REFERENCE: 07180/004003
                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR ETLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
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CURRENT FILING DATE: 2001-04-05
                                                                                                                                                                                                                                       SOFTWARE:
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                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                        LENGTH: 1156
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                                                              SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                                                   CTGGAGGAGCAGGCCCAGCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCCCGCCTCAAG
                                                                                           LeuGluGluAlaGlnAlaGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
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APPLICANT: ZADNIS, Vassilis
APPLICANT: Kypreos, Kyrlakos E.
TITLE OF INVENTION: Compounds and methods fc
TITLE OF INVENTION: Cholesterol levels wit
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827.854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
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; ORGANISM: Homo
US-09-827-854-10
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; OTHER INFORMATION:
US-09-870-759-129
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CURRENT FILING DATE: 2002-01-14
PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 166
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 129
LENGTH: 1156
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 129, Application US/09870759 Patent No. US20020177551A1
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TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 870759
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APPLICANT: Braun, Andreas
APPLICANT: Bonsal Aruna
APPLICANT: Kleyn Patrick
TITLE OF INVENTION: GENES AND POLYMORPHISMS AS:
TITLE OF INVENTION: CARDIOVASCULAR DISEASE AN
FILE REFERENCE: 24736-2048
CURRENT APPLICATION NUMBER: US/09/802,640
CURRENT FILING DATE: 2001-03-09
; NAME/KEY: CDS
; LOCATION: (61)...(1014)
; OTHER INFORMATION: Nucleotide
; OTHER INFORMATION: (APOE)
US-09-802-640-17
                                                                                                                                                                                                                                                                                                                                 RESULT 4
US-09-802-640-17
                                                                                                                                                                                                                                                                                    Sequence 17, Application US/09802640 Publication No. US20030036057A1 GENERAL INFORMATION:
                                                                                                                SEQ ID NO 17
LENGTH: 1156
                                                                                  LENGTH: 1156
TYPE: DNA
ORGANISM: Homo
                                                                                                                                           SOFTWARE:
                                                                                                                                                          NUMBER OF SEQ
                                                                      FEATURE:
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RESULT 5
US-09-827-854-8
Sequence 8, Application US/09827854
Patent No. US20020123093A1
GENERAL INFORMATION:
APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Cholesterol levels with FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US/09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US/09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEO ID NOS: 19
SOCIUM OS
SEQ ID NO 8
LENGTH: 1156
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Best Local Similarity:
Query Match:
DB:
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ORGANISM: Homo sapiens
FEATURE:
NAME_KEY: misc_feature
OTHER INFORMATION: Incyte II
NAME_KEY: unsure
LOCATION: 46
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 454, Application US/10044090 Patent No. US20020137081A1 GENERAL INFORMATION
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriak
                                                                                                                                                                            Sequence 12, Application Patent No. US20020123093
                 SEQ ID NO 12
LENGTH: 1156
ORGANISM: Homo
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
FILE REFERENCE: O7180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SECTION OF SEQ ID NOS: 19
SECTION OF SEQ ID NOS: 19
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                                                                  SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
                                                                                   GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
                                                            AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCGAAG
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Coholesterol levels without inducing hyp
ETILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 1156
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US-09-827-854-11
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US-09-827-854-11
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                  LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
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RESULT 10 US-09-880-107-2244 US-09-880-107-2244 ; Sequence 2244, Application US/09880107 ; Patent No. US20020142981A1 ; GENERAL INFORMATION:	Qy 301 ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 317	Qy 281 SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300	Oy 261 LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 280	Qy 241 SerargThrargAspargLeuAspGluValLysGluGlnValAlaGluValArgAlaLys 260 	Qy 221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240	Qy 201 ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220 	Qy 181 GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200 	Qy 161 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr 180 	Qy 141 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160 	Qy 121 ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140	Qy 101 ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120	Qy 81 LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu 100 	Qy 61 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80	Qy 41 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 60 	Qy 21 GluGlnAlaValGluThrGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40	Qy 1 MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20 	DB: 10 Gaps: 0 US-09-827-854-16 (1-317) x US-09-954-456-760 (1-1157)

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; OTHER INFORMATION: Genbank Accession
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SEQ ID NO 2244
LENGTH: 1157
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PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
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TITLE OF INVENTION: Gene Expression Profiles
FILE REFERENCE: 44921-528-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
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                                GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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Mismatches:
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Percent Similarity:
Best Local Similarity:
Query Match:
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Pred. No.:
Score:
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Sequence 7, Application US/09827854
Patent No. US20020123093A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
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SEQ ID NO 7
FRAGTH: 1156
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without inducing hypertriglyceridemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 07180/004003
                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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                                                                                                                                GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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              LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                 GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
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                                                        GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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Sequence 5, Application US/09967013
Patent No. US20020045840A1
GENERAL INFORMATION:
APPLICANT: Stanton, Jr. Vincent P.
TITLE OF INVENTION: METHOD FOR GENETIC ANALYSIS OF TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: 11926-022001
CURRENT APPLICATION NUMBER: US/09/967,013
CURRENT FILING DATE: 2000-10-25
PRIOR APPLICATION NUMBER: 60/206/613
                                                    ; TYPE: DNA
; ORGANISM: HOMO
US-09-967-013-5
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US-09-967-013-5
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  Percent Similarity:
                                                                                           NUMBER OF SEQ ID NOS: 91
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 5
                                                                                                                         PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 2000-05-23
                                                                                  Q ID NO 5
LENGTH: 41907
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  4.21e-115
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Matches:
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March:	38 127	aGlnAlaArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgG 	11 121	Qy Db	
MATCH::::::::::::::::::::::::::::::::::::	18 121	uGlnLeuThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaA 	9 115	Qу Db	
Match:	115	-ArgAlaLeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluG 	7 109	Qу ДЬ	
Natch   10.10%   Mismatches   194	109	CCCCATCCCAGCCCTTCTCCCCGCCTCCCACTGTGCGACACCCTCCCGCCCTCTCGGCC	103	Db	
Match: 10.2% Indexces: 194  B27-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProGlu 11			78	Qy	
Match:   10   104   Matshacches:   194	103	GCCTCTGCCCGTTCCTTCTCTCCCTCTTGGGTCTCTCTGGCTCATCCCCATCTCGCCC	097	Db	
Match:				Qy	
Match: 80.00% Mismatches: 194  ###################################	097	TCGTCTCTGCCTCTGCCTCTGCATCTGCTCTCTGCATCTGTCTCTGTCTCTCTC	091	Db	
MATCH: 10.00% MISMATCHES: 194  MATCH: 10.12% Indels: 194  10.12% Indels: 194  12. PheLeuAlaGlyCysGlnAlaLysValGluGhnAlaValGluThrGluProGluPr				Qy	
Match: 10.0% Mismatches: 194  Match: 10.0% Mismatches: 194  10.28 Indels: 194  12.27-854-16 (1-317) x US-09-967-013-5 (1-41907)  12.2 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluTrGluProGl	091	CCTCCCAAAGTGCTGGGATTAGAGGCATGAGCCACCTTGCCCGGCCTCCTAGCTCCTTC	280	Дb	
Match: 10.0% Mismatches: 104  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluGluGluGluGluGluGluGluGluGluGluGluGluG				Qy	
Match: 10.00% Mismatches: 194  Match: 10.00% Mismatches: 194  10.000 Gaps: 194  11.000 Gaps: 194  12.PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluTroGluProGlu 31  11.000 Hill Hill Hill Hill Hill Hill Hill H	085	CTCACTGTGTTGCCCAGGCTGGTCTTGAACTTCTGGGCTCAAGCGATCCTCCCGCCTCG	079	Db	
Match: 86.12% Mismatches: b Gaps: 194  10 Gaps: 194  11 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGlaGCCCAGCCCGAGCCCGAGCCCGAGCCCGAGCCCGAGCCCGAGCCCGAGCCCGAGCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCGAGCCCCTGGCAACTGTGCGCAACTGTGCGCAACTGTGCGCAACTGTGCGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGC				Qy	
Match: Similarity: 60.00% Index: 194  Batch: 10 Gaps: 194  B27-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProGlu 31	079	CGTCCTGGCTCTGTCTCTGTCCTTCCCTAGCTCTTTTATATAGAGACAGAGAGATGGGG	073	dd	
Match: Similarity: 60.00% Index: 194  Match: 10 Gaps: 194  10 Gaps: 194  11 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluFroGluGluFroGl				Qy	
Match: 81milarity: 60.00% index: 194  Match: 10  80.12% index: 194  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluglnAlaValGluThrGluProGluGluProGluGluProGluGluProGluProGluGluFillIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	073	TGGAATTCTCTCTCAGCTTTGTCTCTCTCTCTTCCCTTCTGACTCAGTCTCTCACAC	067	Db	
Match: Similarity: 60.00% Index: 194  Match: 10 Gaps: 194  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluFhrGluFroGrogroGroGrogroGrogroGrogroGroGrogroGrogroGrogroGrogroGrogroGrogroGrogroG				Qy	
Match: 80.10% Mismatches: b Gaps: 194  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGlu	067	CCTGGGTCTCTGCTTGTAGCTTCCTCTTCCCATTTCTGACTCCTGGCTTTAGCTCT	061	Db	-
Match: 80.10% Mismatches: b Match: 10  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluThrGluTroGlu				Qy	
Match: Similarity: 60.00% Index: 6 Match: 10 Gaps: 194  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluFillIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	061	GGCGGCTATACCTCCCCAGGTCCAGGTTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGG	055	Db	
Match: Similarity: 60.00% Mismatches: 6 Match: 10 Gaps: 194				Qy	
Match: 86.128 Indels: 194  Match: 10 Gaps: 194  10 Gaps: 1  10 Gaps: 1  10 Indels: 194  10 Gaps: 1  10 Indels: 194  10 Gaps: 1  10 Indels: 194  10 Indels: 194  10 Indels: 194  10 Indels: 194  11 Indels: 194  11 Indels: 194  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGloGLUProGluProGluProGluProGluProGloGLUProGluProGluProGluFlUIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	8	SerGlnValThrGlnGluLeu	7	ФФ	
Match: 80.10% MISMATCHES: 6  Match: 86.12% Indels: 194  827-854-16 (1-317) x US-09-967-013-5 (1-41907)  12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProGlu 31	1		5 043	Дy	
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Match: 86.12% Indels: 19 Match: 10 Gaps: 1 827-854-16 (1-317) x US-09-967-013-5 (1-41907)	1 037	PheLeu     TTCCAC	1 031	Qу Дъ	
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APPLICANT: NOSEN et al.
APPLICANT: ROSEN et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and FILE REFERENCE: PA104
CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT//S00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 50/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 896
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 133
LENGTH: 786
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Best Local Similarity:
Query Match:
DB:
                                                                                                             US-09-827-854-16 (1-317) x US-09-925-302-133 (1-786)
                                                                                                                                                                                                 Score:
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                                                      CGCCTGGTGCAGTACCGCGGCGAGGTGCASGCCATGCTCGGCCAGAGCACCGAGGAGCTG
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Best Local Similarity:
                                            US-09-827-854-16 (1-317) x US-09-964-824A-374 (1-478)
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CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
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Patent No. US20020102531A1
GENERAL INFORMATION:
APPLICANT: Horrigan, Stephen
                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.0 SEQ ID NO 374 LENGTH: 478
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign TITLE OF INVENTION: Sets FILE REFERENCE: 689290-73
                                                                                                                                                                                                 NAME/KEY: misc_feature LOCATION: (1)...(478) OTHER INFORMATION: n=a.
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APPLICANT: Horne, Darci T.

APPLICANT: Vockley, Joseph G.

APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.

TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/231,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2491
ENCOTH: 478
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US-09-880-107-2491/c
; Sequence 2491, Application
; Patent No. US20020142981A1
; GENERAL INFORMATION:
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Best Local Similarity:
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ORGANISM: Homo sapiens
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Command line parameters:

-MODEL-frame+_p2n.model -DEV=xlp
-O_-Ggn2_1/USPTO_Spool_US09827854/runat_11032003_101610_27495/app_query.fasta_1.3576
-DB=EST -OFMT-fastap -SUFFIX=p2n.rst -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT=0
-UNITS-bits -STARF-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN-0 -MAXLEN-200000000
-USER-US09827854_8CGN_1_1_8826_@runat_11032003_101610_27495 -NCPU-6 -ICPU-3
-NO_SLPXY -NO_MMAP -LARGEQUERY -NEC_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120
-WARN_TIMEOUT-30 -THREADS-1 -XGAPOD+10 -XGAPEXT-0 -S -FGAPEXT-7
-YGAPPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

## ALIGNMENTS

AUTHORS TITLE	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	в1670367	RESULT 1
NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 1027)	Homo sapiens	human .	EST.	BI670367.1 GI:15584600	B1670367	mRNA sequence.	603292738F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5312024 5',	B1670367 1027 bp mRNA linear EST 12-SEP-2001		

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI),
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished
Contact: Rob
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High quality sequence stop: 845.
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/db_xref="taxon:9606"
/clone="IMAGE:5312024"
/clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note-*Organ: brain; Vector: pBluescriptk (modified pBluescript KS+); Site_1: BamHI; Site_2: Sall-Xhol (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, NATIONAL Institutes of Health). Note: this is a NIH_MGC Library." a 308 c 400 g 125 t
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                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
Tissue Procurement: ATCC/DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llni.gov
Plate: LLCM1718 row: b column: 10
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 842)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BG763371 842 bp mkNA lined For For CONA clone IMAGE:4860585
                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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BG763371
                                                                                                                                         quality sequence stop: 817.
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/clone="IMAGE:4860585"
/clone_lib="NIH_MGC_496"
/tissue_type="melanotic melanoma, high MDR (cell line)"
/lab_host="DH10B (phage-resistant)"
                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                        1. .842
                                                                                                                     Location/Qualifiers
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DB: Query

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Alignment Pred. No.: Score:

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                                                                                                                                                                                   CAGCCGGACCCGCGACCGCTGGACGAGGTGAAGGGACCAGGTGGCGGAGGTGCGCGCAAG
                 ySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
                                                                                                                       GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                                                                                                                                                                                                                                                                                                            ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
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                                                             CTACAGGAGCGGCCCAGGCCTGGGGCGAGCGGCGCGCGGGATGGAGGAGATGCGG
                                                                          LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMet-Gl
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ECCRI; CDNA made by oligo-dT priming. Directionally cloned
into ECCRI/XhoI sites using the following 5' adaptor:
GGCACGAG(G). $ize-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-CDNA synthesis kit (Stratagene) and Superscript
ILRT (Life Technologies). Note: this is a NIH_MGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Library. |"
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GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40
                                                           GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCAGACCGAGTGGCAGAGC
                                                                                                                                         ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tissue Procurement: Miklos Palkovits, M.D., Ph.D.

CDNA Library Preparation: Michael J. Brownstein (NHGRI), St
Toshiyuki and Piero Carninci (RIKEN)

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999)
Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 93)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
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Mammalia; Eutheria; Primates;
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                                                                                                                                             mRNA sequence.
BG761746
BG761746.1 GI
EST.
Unpublished (1999)
Contact: Robert Strausberg, ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium ()
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae 1 (bases 1 to 938)
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/Clone="IMAGE:4841411"
/Clone=lib="NHI_MGC_49"
/Clone=lib="NHI_MGC_49"
/tissue_type="melanotic melanoma, high MDR (cell line)"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/lab_host="Organ: skin; Vector: poTB7; Site_1: xhoI; Site_2:
/note="Organ: skin; Vector: poTB7; Site_1: xhoI; Site_2: line of Site_3: skin; Vector: poTB7; Site_3: skin; Vecto
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11762 row: a column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 942) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)
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603249241F1 NIH_MGC_96
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Location/Qualifiers
                                                                                         /clone_lib="NiH_MGC_96"
/clone_lib="NiH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
/lab_host="DH10B"
/lab_host="DY0gan: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: Sal1-xhoI (gtcgag.); Oligo-dT primed using primer 5'-TTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carnincl, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."
77 a 284 c 361 g 120 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
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                                                                                                                               GCCTCAAAAC - TGGTTGACCCCT
                                                                                                                                                            rgLeuLysSerTrpPheGluPro
                                                                                                                                                                                                          laLysLeuGluGluGlnAla---GlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaA
                                                                                                                                                                                                                                                       GCAGCCGGACCCGCGACGGCCTGGACCGAGGTGAACGGACCAGGTTGCGGAAGTTGCGCG
                                                                                                                                                                                                                                                                                                                      CCCCTGGTGGAACAGGGCCGCGTGCGGGC - GCCACTGTGGGCTCCCTGGCCGGCCAGCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CAGGCCGGGGCCCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
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mRNA sequence.
BM042094
BM042094.1 GI
                                              BM042094
603615713F1
   GI:16771361
                                              NIH_MGC_112
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Indels:
                                              bp mRNA linear EST 07-NO sapiens cDNA clone IMAGE:5420618
                                                               mRNA
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EST 07-NOV-2001 E:5420618 5',

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Best Local Similari
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ORIGIN
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                                                                                   LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluGlnLeu
                                                                                                                                                       LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                                                                                                                      GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                                                                                                                                                                        CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
                                                                                                                                                                                                         GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
ACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
                                                                   CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 800)
11 (http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           http://image.llnl.gov
Plate: LLCM1875 row: i column:
High quality sequence stop: 792.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     found through the I.M.A.G.E. Consortium/LLNL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Seguencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2: ECORI; cDNA made by oligo-dT prining. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
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/clone="IMAGE:5420618"
/clone_iib="NIH_MGC_112"
/clone_iib="NIH_MGC_112"
/tissue_type="melanotic melanoma, cell line"
/lab_host="DH10B_(phage_resistant)"
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1195.00
98.80%
97.20%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MCC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: image.llnl. Plate: LLCM1419 row: k column: 08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          National Institutes of Health, Mammalian Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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BG472299
BG472299.1 GI:13404485
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Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 927)
NIH-MGC http://mgc.nci.nih.gov/.
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                170
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/lab_host="DH10B (phage_resistant)"
/lab_host="DH10B (phage_resistant)
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/lab_host="DH10B (phage_resistant)"
/lab_host="DH10B
                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4645759"
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                                      CCGGCCTCAAGAGCTGGTTCGGAGCCCTGGGTGGAAGACATG
                                                 aArg-LeuLysSerTrpPhe-GluProLeuValGluAspMet
                                                                                      gAlaLysLeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAla---PheGlnAl
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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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BI597743
BI597743.1 GI
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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Location/Qualifiers
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/db_xref="taxon:9606"
/clone="IMAGE:5300259"
/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 817)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
                                                                                            CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLCM1612 row: e column: 20
                                                                                                                                                                   Contact: Robert Strausberg, I
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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602649975F1 NIH_MGC_40
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Location/Qualifiers
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4760827"
/clone_lib="NIH_MGC_40"
/tissue_type="carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
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                  LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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CTGTCTGAGCAGGTGCAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
                                                                      GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11726 row: g column: 10
High quality sequence stop: 783.
Location/Qualifiers
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (Dases 1 to 811)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
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/db_xref="taxon:9606"
/clone="IMAGE:5287329"
/clone_lib="NIH_MGC_96"
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                                                                                                                                cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1790 row: g column: 17 High quality sequence stop: 829.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 845)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mRNA sequence.
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                                                                                                                       1. .845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GI:14177059
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   poTB7; Site_1: XhoI;
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                                                                                                               ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
                                                                                                                                                                  GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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           SerArgThr-ArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
                                                 CTACAGGAGCGGGCCAGGCTG---GGGCGAGCGGTGCGCGCGCGGATGGAGGAGATGGGC
                                                                                                   CCCCTGGTGGAACAGGGCCGCGTGCGGGCCGC-ACTGTGGGCTCCCTTGGCGGCAG-CCG
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Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GCAACGAG(G), Size-selected >500bp for average insert size 1.8kb. Library constructed by Li Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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TITLE
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Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLAM10672 row: e column: 21
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BG707147.1 GI:13983201
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National Institutes of Health, Mammalian
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consorti
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BI551475
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603194314F1 NIH_MGC_95
                                                                                        DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11691 row: 1 column: 04
High quality sequence stop: 812.
                                                                                                                                                                                                                     Unpublished (1999)
Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                     Mammalia; Eutheria; Primates; Catarrhini; 1 (bases 1 to 919)
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National Institutes of Health,
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/clone_lib="NIH_MGC_95"
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                                                        /organism="Homo
                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                           Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Homo :
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Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: L/CMD021 row: k column: 21 High quality sequence stop: 757.
Location/Qualifiers
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Tissue Procurement: DCTD/DTP
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National Institutes of Health, Mammalian
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/lab_host="PHIOB (phage resistant)"
/note="Organ: skin; Vector: pOTBP; Site_1: XhoI; Site_2:
/note="Organ: skin; Vector: poTBP; Site_1: XhoI; Site_2: 
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                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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National Institutes of Health, Mammalian
Unpublished (1999)
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602669093F1 NIH_MGC_96
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                                                                                                                                                                found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
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Command line parameters:

-MODEL-frame+_p2n.model -DEV=xlp
-O_cgn2_1/GSPTO_Spool_VGS09827854/runat_11032003_101610_27486/app_query.fasta_1.3576
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-VGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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AOLK67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC
C12N15/09//C07K14/775,
PC (C12N5/10, C12R1:91), C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
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Patent: JP 1985118189-A 1 2
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PN JP 1985118189-A/1
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PN JP 1985118189-A/1
PI TERANISHI YUTAKA, TAKA
KIMURA MASAKO,
PI IKEDA YASUKO
PC C1XN15/00,C07H21/04//C
CC strandedness: Double;
CC topology: Linear;
CC anti-sense: No;
CC anti-s
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Teranishi,Y., Takamatsu,N., Matsui,Y., Kimura,M. and DNA FRAGMENT
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E00359.1 GI:21686
JP 1985118189-A/1.
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25-JUN-1985
29-NOV-1983 JP 1983224980
TERANISHI YUTAKA, TAKAMATSU
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C12N15/00,C07H21/04//C12P21/00;
strandedness: Double;
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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S Teranishi,Y., Matsui,Y., Ikeda,Y. and Kimura,M.

S Teranishi,Y., Matsui,Y., Ikeda,Y. and Kimura,M.

S Teranishi,Y., Matsui,Y., Ikeda,Y. and Kimura,M.

Patent: JP 1986096997-A 1 15-MAY-1986;

MITSUBISHI CHEM IND LTD

OS Human (Homo sapiens)

PN JP 1986096997-A/1

PD 15-MAY-1986

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

PF C C12P1/00,A61X35/74,A61X37/04,C12N15/00,(C12P)

PC C12R1:19);

CC (c12N15/00,

PC C12R1:19);

CC strandedness: Double;

CC topology: Linear;

CC hypothetical: NO;

CC *source: tissue_type=Liver;

FH Key

FF Sig_peptide 15. 68

FT Sig_peptide 15. 68

FF Sig_peptide 15. 68
ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
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TERANISHI YUTAKA, MATSUI YASUSHI, IKEDA YASUKO, KIMURA MASAKO C12P21/00, A61K35/74, A61K37/04, C12N15/00, (C12P21/00, C12R1:19),
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/db_xref="taxon:32644"
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                         Tumor markers in ovarian cancer Patent: WO 0175177-A 63 11-OCT-2001 THE SECRETARY OF THE DEPARTMENT OF
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/organism="Homo
/db_xref="taxon:
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Mammalia; Butheria; Primates; Catarrhini;

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1 (bases 1 to 1156)

Fujita,S., Hamanaka,H., Fukui,Y. and Yoko

Apo E humanized mammal

Patent: JP 2001017028-A 2 23-JAN-2001;

MITSUBISHI CHEMICAL CORP

OS Homo sapiens (human)

PN JP 2001017028-A/2

PD 23-JAN-2001

PR 31NOBU FUJITA,HIRKI HAMANAKA,YUKO

PR SHINOBU FUJITA,HIRKI HAMANAKA,YUKO
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A01K67/027,A61K45/00,A61P25/28,A61P43/00,C12N5/10, PC
C12N15/09//C07K14/775,
PC (C12N5/10,C12R1:91),C12N5/00,C12N15/00,(C12N5/00,C12R1:91)
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/db_xref="taxon:9606"
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Breslow, J.L.
Synthesis, intra
apolipoprotein 1
                                                                                                                                                                                                                                                                                                                                                                                       HUMAPOE3
                                                                                        Williamson,R. and Humphries,S. The isolation of cDNA clones for detection of apoE RNA in hepatic EMBO J. 2 (12), 2369-2373 (1983)
                                                                                                                                                                                                                          Breslow,J.L., McPherson,J., Nussbaum,A.L., Williams,H.W., Lofquist-Kahl,F., Karathanasis,S.K. and Zannis,V.I. Identification and DNA sequence of a human apolipoprotein
                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 355 to 1156)
                                        Zannis, V.I.,
                                                                                                                                               Wallis, S.C.,
                                                                                                                                                                         6897404
                                                                                                                                                                                       J. Biol. Chem.
83082756
                                                                                                                                                                                                               clone
                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                   apolipoprotein; apolipoprotein low density lipoprotein.
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K00396.1 GI:178850
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preapolipoprotein E
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                                                                                                                                                                                                  (24),
             processing,
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                                      Goldberger, G.,
                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Catarrhini; Hominidae
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                                                                                                                                               Markham,A.,
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(APOE)
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             signal
                                        Karathanasis, S.K. and
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WA, complete
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Biol. Chem. 258, 11422-11422 (1983)] correction of [1]. [4] epsilon-3 and variant.
[5] epsilon-2 aliele.
[6] sites; mntation-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Apo E is a component of normal human very low density lipoproteinemia. There are six human apo E phenotypes known to result from a single structural gene, three of the common alleles being epsilon-4, epsilon-3 and epsilon-2. This sequence appears to be of the epsilon-3 allele. [1] argues that the apo E polymorphism involves mutations in the structural coding region; for example the epsilon-2 phenotype which is characterized by hyperlipoproteinemia is thought to result from a c to t change (arg to cys) at base 586 below [3],[5]. The sequence shown is 57% homologous with human apo A-I and 81% homologous with rat apo E. For the epsilon-4 sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  J. Biol.
84185684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Apo E is located on chromosome 19 --Jackson, Bruns and Breslow , PNAS USA 81, 2945-2949 (1984) -- and is believed to be linked to the Apo C-II gene (see separate entry).

The two mutations causing type III hyperlipoproteinemia (apollpoprotein E phenotype E3/3) produces substitutions of Arg for (apollpoprotein E phenotype E3/3) produces substitutions of Arg for Cys at amino acid 112 and Cys for Arg at amino acid position 142.

Draft entry and printed copy of sequence for [1] kindly provided by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Humphries, S.E., Cumming, A.M. and Hardman, N.
Isolation and characterisation of a variant allele of the human apolipoprotein E
Biochem with the state of the prochem because of the pr
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J. Biol. Chem. 259 (10), 6498-6504 (1984)
84212473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              L.L. Gill, University of Aberdeen. Complete source information:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              See the separate entry.

[J. Biol. Chem. 258, .11422-11422 (1983)] is too extensively revised relative to [1] to record the revisions in the FEATURES table. The sequence below is that of [J. Biol. Chem. 258, 11422-11422 (1983)]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J. Clin. Invest. 83 (4), 1095-1101 (1989)
89198059
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   85279526
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Gill, L.L., Peoples, O.P.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 McLean, J.W.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                               [4] and blood [5], cDNA to mRNA.
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/protein_id="AAB59546.1"
                                                                                                    /gene="APOE"
                                                                                                                                                                                                                                                                        /map="19q13.2"
                                                                                                                                                                      /gene="APOE"
                                                                                                                                                                                                                                  tissue_type="liver"
                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                            location/Qualifiers
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note="g in
                                  'gene="APOE"
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/citation=[2]
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369
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288
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/citation=[4]
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/citation=[4]
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                                                                                                                                                    GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                             ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
                                                                                                                                                                                                                                                                   ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140
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SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                                                   AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
                                                                                                                                                                                               LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
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BC003557
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Submitted (20-FEB-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer General Cance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clone distribution: MGC clone distribution through the I.M.A.G.E. Consortium/LLNL at: Series: IRAL Plate: 6 Row: h Column: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy, George Yang, Scott Zuyderduyn, Marco Marra.
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BC Cancer Agency, Vancouver, BC, Canada
info@bcgsc.bc.ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Genome Sequence Centre,
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Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Mammalia; Eutheria; Primates;
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QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
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/protein_id="AAH03557.1"
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/translation="MKVLWAALLVTFLAGCQAKVEQAVETEPEPELRQOTEWQSGQRW
ELALGRFWDYLRWVQTLSEQVQEELLSSQVTQELRALMDETWKELKAYKSELEEQLTP
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/clone_lib="NIH_MGC_16"
/lab_host="DH10B-R"
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/clone="MGC:1571 IMAGE:3355712"
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/db_xref="LocusID:348"
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      ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
                        SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                  LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
                                                                                                                                                      GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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                                                                           AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCCAAG
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METHOD FOR PRODUCING APPLIPOPROTEIN
Patent: JP 1994315392-A 1 15-NOV-1994;
MITSUBISHI KASEI CORP
OS HOMO SAPIENS (human)
PN JP 1994315392-A/1
PD 15-NOV-1994
PF 11-JUN-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF (12P21/02,CO7K13/00,C12N5/10,C12N15/PC C12R1:91);
PC (C12R1:91)
CC strandedness: Double;
CC (C12N5/10,C12R1:91);
CC strandedness: Double;
CC topology: Linear;
FH Key Location/Qualifiers
FH Key Corganism='Homo sapier Sig_peptide 15..68
FT sig_peptide 59..968
FT mat_peptide 69..968
FT mat_peptide 69..968
                                   GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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JP 1994315392-A/1.
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JP 1994315392-A/1
15-NOV-1994
11-JUN-1985 JP 1994015433
MORIMOTO HIRONORI, TERANISHI YUTAKA
C12P21/02,C07K13/00,C12N5/10,C12N15/12,C12N15/18,(C12P21/02,
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/db_xref="taxon:9606"
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Location/Qualifiers
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Sequence 378
AX333278
AX3333278.1
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Horrigan, S., Soppet, D.R. and
Cancer gene determination and
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Mammalia; E
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                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
370 c 426 q 14
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2.69e-83
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Patent W00194629.
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Primates;
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Length:
Matches:
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Mismatches:
Indels:
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                                   CAGGCCGGGGCCCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
                                                                             LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
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/db_xref="taxon:9606"
/note="EMBL/GenBank Acce.
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Attie,A.D., Gretch,D.G., Sturley,S.L.
Production of recombinant proteins in
Patent: US 5472858-A 1 05-DEC-1995;
Location/Qualifiers
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1. 1157
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1. .1157
                                                              /clone="pHAE[112,178,813]."
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KLRKRLLRDPDDLQKRLAYVQAGAREGAERLSALTERELGPLVEQGRVRAATVGSLAG
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QPLQERAQAWGERLRAKMEEMGSTRORLDEVKEQVAEVFAKLEEQAQQIRLQAEAFQ
ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH"
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/note="apoE mRNA;
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/replace="c"
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Patent: JP 2001017028-A 3 2:
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/3
PN JP 2001017028-A/3
PD 23-JAN-2001
PF 28-APR-2000 JP 20001289
PR SHINOBU FUJITA, HIROKI I
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JP 2001017028-A/3.
        PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE YOKOYAMA
AOIK67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC
C12N15/09/C07K14/775,
PC (C12N5/10,C12R1:91),C12N5/00,C12N15/00,(C12N5/00,C12R1:91)
                                                                                                                         Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                              AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCGCCAAG
                                       SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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X13887 GI:38054
apolipoprotein; apolipoprotein E Macaca fascicularis.
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Direct Submission
Submitted (16-DEC-1988) Marotti K.R., The
301 Henrietta Street, Kalamazoo, MI 49008
2 (bases 1 to 1178)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Data
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/note="polyA signal"
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/clone_lib="lambda gt
83. .1036
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/db_xref="GI:3|8055"
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Location/Qualifiers
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Eukaryota; Met
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Direct Submission
Submitted (27-APR-2000) Department of Molecular Biotechnology,
Submitted (27-APR-2000) Department of Molecular Biotechno
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Nickerson,D.A., Taylor,S.L., Fullerton,S.M., Weiss,K.M., Clark,A.G., Stengard,J.H., Salomaa,V., Boerwinkle,E. and Sing,C.F. Sequence diversity and large-scale typing of SNPs in the human apolipoprotein E gene Genome Res. 10 (10), 1532-1545 (2000)
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Nickerson, D.A.
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/map="19q13.2"
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/replace="t"
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/note="putative"
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/replace="t"
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308
/codon_start=1
                             join(1878. .1920,3013.
/note="APOE"
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join(1060. .1094,1855. .1920,3013. .3205,3786. .4645)
product="apolipoprotein-E"
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/note="putative"
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/rpt_family="MIR"
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                  frequency="0.45"
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/note="putative"
/rpt_family="Alu"
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/replace=""
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ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH"
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-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Qualifiers		apoE; cholesterol; atheroscleros w density lipoprotein; LDL; ds.	) allele, apoEl DNA.			BP.	ALIGNMENTS	AAH98479 AAA44191	AAC02139 AAC011980	ABL6/340 AAS03049	ABL62679	ABK34238 ABN95994	AAF18114 AAV89595	AAS22673	AAT18068 AAO69099	AAT18070 AAT69792	AAQ69101	AAD32075 AAV29159	AAD32077	AAD26108	AAZ09524 AAZ09526	AAD26034	AAX75756	AAD22047 AAS22437	AAF84316	ABL65450	ABN95746	AAT06957	AAD22052	AAF84315 AAD22048	ABA83113	260	D220	AAD22050	ID
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                                                                                                                                                                                                                                                                                                                                                                                                      New apolipoprotein E polypeptide and nucleic acid, useful for l cholesterol, delaying the onset of or treating atherosclerosis mammal, without inducing hypertriglyceridemia -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New apoE humanized mammalian cell useful for treating or preventing arteriosclerosis -
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LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
                                   ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
                                                            CAGGCCGGGGCCCGAGGGCGCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
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                          AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGTGCCTGGCAGTGTAC
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)B; AAB80996.
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                                                        New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia -
                                             Claim
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04-OCT-2000; 2000US-06790B8
05-APR-2001; 2001US-0827854
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hypertriglyceridaemia; low density lipoprotein; LDL; ds.
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The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE2 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atheroselerosis, treating or regressing atheroselerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (L

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 The patent dis E (APOE) gene.
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                                                                                       Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene -
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                                                       Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for haplotyping and/or genotyping the APOE gene. The haplotyping methods of the invention are useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with APOE activity, e.g. familial dysbetalipoproteinaemia, type III hyperilipoproteinaemia, atherosclerosis, and Alzheimer's disease. They are useful to validate APOE as a candidate agent for treating a specific condition or disease predicted to be associated with APOE activity and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with APOE activity. Genotyping or haplotyping methods are useful to screen for compounds targeting APOE to treat a specific condition or disease associated with APOE activity. The present secures is a none associated with APOE activity.
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                                                                                                      GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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                                                                                          CAGGCCGGGGCCCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
                                                                                                                                                         LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
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                                        AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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                                                                             The encoded protein is used to treat subjects who are deficient in apolipoprotein-E (or who produce abnormal forms of this molecule) and therefore are likely to suffer from hyperlipidemia, resulting arteriosclerosis. It can also be used to raise antisera for detecting the protein deficiency or production of abnormal forms.
                                                               Sequence 1110
                                                                                                                              Disclosure; Fig 2; 45pp; English
                                                                                                                                              New DNA sequence coding for human apolipoprotein-E expression vectors and transformed cells contg. it
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MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal

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(1-1110)

US-09-827-854-17 (1-317) x AAN60409

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CC The invention relates to methods for diagnosing and prognosing ovarian CC tumours in an individual via the detection and measurement of the CC expression of ovarian tumour marker genes (ABA83081-ABA83122, ABA83180, CC ABA83182 and ABA83184) or segments thereof (ABA83123-ABA83122, ABA83179, CC ABA83181 and ABA83183). The methods of the invention are useful for detecting an ovarian tumour in a patient, for identifying an individual cassessing the relative severity of ovarian cancer, in prognostic tests for monitoring a patient in remission from ovarian cancer and in tests for monitoring a patient in remission from ovarian cancer and in tests for CC monitoring a patient in remission from ovarian tumour selected from cc. The methods can additionally be used to identify a particular tumour as CC being an ovarian tumour (i.e., an epithelial ovarian tumour selected from CC serous cystadenoma, borderline serous tumour, serous cystadenocarcinoma, condentation mucinous tumour, mucinous cystadenocarcinoma, condentation of constance in indifferentiated carcinoma, CC clear cell adenocarcinoma, cystadenofibroma, adenofibroma and Brenner CC tumour. The ovarian tumour marker genes of the invention were identified coverexpressed in a broad variety of ovarian epithelial tumour cells crelative to normal ovarian epithelial calls. The marker genes are implied to immune response nathways, in the resultation of cell
                                                                 implicated in immune response pathways, in the regulation of cell proliferation and in protein folding, and many of these are membrane-localised or secreted. In addition to their use as diagnostic and prognostic markers, the ovarian tumour marker genes or their encoded proteins may be used as therapeutic targets for the treatment and prevention of ovarian cancer. Sequences ABA83081-ABA83122, ABA83180, ABA83182 and ABA83184 represent the ovarian tumour marker genes of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mucinous cystadenoma; borderline mucinous tumour; endometrioid carcinoma undifferentiated carcinoma; clear cell adenocarcinoma; cystadenofibroma; adenofibroma; brenner tumour; serial analysis of gene expression; SAGE; immune response pathway; cell proliferation regulation; protein folding; membrane localised; secreted; therapeutic target; cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting and identifying ovarian tumor, identifying increased risk for developing ovarian cancer, and determining effectiveness of ovarian cancer treatment, by measuring expression level of ovarian tumor marker
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P-PSDB; ABB50287.
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                                                                                                                                                                                                                                              Sequence 1156
                                                                                                                                                                                                                                                                                                      Disclosure; Page 13-14;
                                                                                                                                                                                                                                                                                                                    useful for treating arteriosclerosis -
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P-PSDB; AAB80997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (Li receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New apolipoprotein E polypeptide and nucleic ac cholesterol, delaying the onset of or treating mammal, without inducing hypertriglyceridemia
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LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
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New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia hypertriglyceridemia

91pp; English.

The present sequence is a human apolipoprotein E (apoE) allele, apoE2\*\* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atheroscierosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LE receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or had alled the content of the protein remnants of the bloodstream or had alled the content of the protein remnants of the bloodstream or had alled the content of the protein remnants of the bloodstream or had alled the content of the protein remnants of the bloodstream or had alled the content of the protein remnants of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the content of the bloodstream or had been supported by the bloodstr or having (LDL)

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Length:
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US-09-827-854-17 (1-317) x AAD22052 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTTGGGATTACCTGCGCTGGGTGCAGACA GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCAGAGCCGAGTGGCAGAGC GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAGGTCACCCAGGAACTGAGGGCG (1-1156)100 420 120 80 60 180 40 120 360 300 240 20

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04-OCT-2000;
05-APR-2001;
             The present sequence is a human apolipoprotein E (apoE) allele, apoE2* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing
                                                                                                               New apolipoprotein E polypeptide and nucleic acid, useful for cholesterol, delaying the onset of or treating atherosclerosis mammal, without inducing hypertriglyceridemia -
                                                                                       Claim
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                                                                                                                                                                                      Recombinant human apolipoprotein-E (ApoE) (AAR86791) can be produced by preparing a genetic construct (contg. an ApoE-encoding sequence, e.g. AAT06957, and flanking regulatory sequences enabling the protein to be expressed in insect cells), which is then introduced into a Manduca sexta larva (using a récombinant Autographica californica nuclear polyhedrosis virus) and recovering the protein from the haemolymph of the larval host. The ApoE produced is in a form sufficiently complexed with lipids to be biologically active, which cannot be achieved in insect cell cultures, and can therefore be
                                                                                                                                                                                                                                                                                                                                                                                  Prodn. of recombinant apo:lipoprotein E in insects - by infecting Manduca sexta larvae with recombinant Autographica californica nuclear polyhedrosis baculovirus vector.
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CC Note: The sequence data for this patient did not form part of the printed consistency in the printed consistency in the progression profiles who but was obtained in electronic format directly from WIPO at figure 1. The proposition of the printed consistency in the progression profiles who is the printed consistency in the printed co
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of diagnosing (I) the onset or CC progression of benign prostatic hyperplasia (BPH), or screening (II) for CC or identifying an agent that modulates the onset or progression of BPH. CC The method is based on changes in gene expression in BPH tissue isolated CC from patients exhibiting different clinical states of prostate (CC hyperplasia as compared to normal prostate tissue. (I) comprises CC detecting the expression levels of one or more genes in prostate cells (II) comprises preparing a first gene expression profile prostate cells. (II) comprises preparing a first gene expression profile CC agent, preparing a second gene expression profile of BPH cells or BPH-like cell population, exposing the cells to the CC agent, preparing a second gene expression profiles. (II) is useful for diagnosing the first and second gene expression of BPH. (II) is CC useful for diagnosing the onset or progression of BPH. (II) is CC useful for diagnosing the onset or progression of BPH. (II) is CC useful for diagnosing the onset or progression clevel in a tissue or cells, by comparing the expression clevel in a tissue or cells, by comparing the expression clevel of genes given in the specification in the tissue or cells to the CC level of genes given in the specification in the tissue or cells to the CC expression of gene in the database, and displaying the CC therefore the expression level in BPH. Agents using (II) are useful for the compared to the expression level in BPH. Agents using (II) are useful for CC therefore the prostate cancer. ABK64106-ABK64860 represent human contents to the prostate cancer. ABK64106-ABK64860 represent human contents to the content and cancer to the content human contents to the content and cancer to the content human contents to the content cancer.
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2000US - 234034P.
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                                                                                                                                                                                                                                                                The present invention describes a method (MI) for screening for an CC anti-neoplastic agent. The method involves exposing cells to a chemical cagent to be tested for anti-neoplastic activity, determining a change in CC expression of at least one gene (I) of a signature gene set, where (I) CC comprises a sequence (S) selected from 8447 sequences (given in ABL6164 CC to ABL70110), or is at least 95% identical to (S), where a change in CC expression is indicative of anti-neoplastic activity. (I) has cytostatic CC activity and can be used in gene therapy. MI can be used for screening CC an anti-neoplastic agent, and can be used for producing a product which cC is the data collected with respect to the anti-neoplastic agent as a CC result of MI, and the data is sufficient to convey the chemical CC treatment of cancer such as colon, breast, stomach, lung, thyroid, CC cesophageal, ovarian, kidney, prostate or pancreatic cancer, infiltrating dotal cancer, squamous cell carcinoma, neuroendocrine cC carcinoma, papillary carcinoma and Wilm's tumour.
                                                                                                                                                                                                                                               adenocarting lower infiltrating lower commonway papillary c
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LeumetAspGluThrMetLysGluLe	GlyGlnArgTrpGluLeuAlaLeuGl 	TGA TGA 110 111	854-17 (1-317) x AAN50450	Scores: 3.93e-113 1560.00 milarity: 98.74% Similarity: 98.74% h: 98.17%	nce 1110 BP; 198 A; 355 C;	sequence may be used to prostment of hyperlipaemia.	3; Page 484; 8pp; Japanes	DNA fragment - contg. DNA which treatment of hyperlipaemia.	1985-188003/31. DB; AAP51204.	) MITSUBISHI CHEM IND KK	V-1983; 83JP-0224980.	V-1983; 83JP-0224980.	N-1985.	18189-A.	eptide 69965 /*tag= b	cation 968	sapiens.	encoding human apolic
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Maximum Match 100
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Copyright (c) 1993 - 2003
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/FCTUS_COMB.seq:*
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US-07-849-389-6
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US-09-287-141-24
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                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, v
CURRENT APPLICATION NUMBER: US/07/709,949
FILING DATE: 19910604
CILASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 962969180
                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
  INFORMATION FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Attie, Alan D
APPLICANT: Gretch, Daniel G
APPLICANT: Sturley, Stephen L
APPLICANT: Beckage, Nancy E
TITLE OF INVENTION: Production of
TITLE OF INVENTION: Insect Larvae
                     TELEPHONE: (608) 251-5000
                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Quarles &
                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                     STREET: P.O. Box
CITY: Madison
STATE: WI
COUNTRY: USA
ZIP: 53701-2113
            TELEFAX:
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                                                                                                            US/07/709,949
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US-09-495-444-24
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US-08-933-577-4
PCT-US92-08-34-4
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US-07-853-913-3
US-08-968-751-26
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US-08-968-751-26
US-08-968-751-3
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US-09-183-861-7
US-09-183-861-34
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US-08-195-196-33-24-3
US-08-467-781-3
US-08-467-853-913-1
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                                             9629691801
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Sequence Sequence

SEQUENCE CHARACTERISTICS:
LENGTH: 1157 base pairs

TYPE: NUCLEIC ACID
STRANDEDNESS: double

TOPOLOGY:

linear

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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-827-854-17 (1-317) x US-07-709-949-1 (1-1157)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; RELEVANT RESIDUES US-07-709-949-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match:
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NAME/KEY: misc_signal
LOCATION: 62..119
PUBLICATION INFORMATION:
NUMBER OF THE PROPERTY OF THE PROPER
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ORIGINAL SOURCE:
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JOURNAL: Journal of Biological Chemistry
VOLUME: 259
PAGES: 6498-6504
DATE: 1984
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LOCATION: 62..1015
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                                                                                                                                                                                 GlnAlaMetLeuAspGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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  GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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Patent No. 6271436
GENERAL INFORMATION:
                                                                    NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEPHORE: (512) 418-3000
TELEPHORE (513) 789-2679
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                   APPLICATION NUMBER: US/08/9
FILING DATE: CONCULTENTLY H
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/C
FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 60/C
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: USANE: DAY/A WANEE.
                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                          STATE: 1A
COUNTRY: US
77210-4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Bazer, Fuller TITLE OF INVENTION: Compo TITLE OF INVENTION: Gener
                                                       SEQUENCE CHARACTERISTICS:
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            TYPE: nucleic acid
STRANDEDNESS: single
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TOPOLOGY:
                                        LENGTH:
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Generation of Transgenic
                                                                                                                                                                                                                                                                                                                                   Release #1.0, Version
                                                                                                                                                                                                      US 60/046,094
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CTCCTCAAAGGCTGGTTCGAGCCTCTGGTGGAAGACATACGGCGCCAGTGGGCCGGGCTG
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; Sequence 5, Applicatio
; Patent No. 6369294
; GENERAL INFORMATION:
                                                                                                                                                             Percent Similarity:
Best Local Similarity:
Query Match:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/819,964
FILING DATE: 28-Mar-2001
CT ROSTETTANTO:
                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/046,094 FILING DATE: 09-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 28-Mar-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: ARNOLD, WH
STREET: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 1126 base pairs
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                                                                                                                                                                                                                                                                                                                             FEATURE
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                          GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu
                                                     Scores:
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Hibler, David W.
REGISTRATION NUMBER: 41,0
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/949,155 FILING DATE: <Unknown>
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1123.00
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Matches:
Conservative:
Mismatches:
Indels:
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RESULT 4
US-08-949-155-51
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                                                                                       APPLICANT: Piedrahita, Jorge A
APPLICANT: Bazer, Fuller W
TITLE OF INVENTION: Compositions a
TITLE OF INVENTION: Generation of
NUMBER OF SEQUENCES: 51
                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE AND DURKEE
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CITY: H
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COUNTRY: US
ZIP: 77210-4433
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                                       Houston
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                                                     P.O. Box 4433
                                                                                                      Compositions and Methods
Generation of Transgenic
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Best Local Similarity:
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TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TA
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 800
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 60/027,338
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
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                           TTGGGGGGAAAAACTTTTTTTTTTTTTTCATTTGACTTCATGTCTTGCTTTCCTTTCC
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                                                                                                          Sequence 51, Application US/09819964
Patent No. 6369294
GENERAL INFORMATION:
APPLICANT: Piedrahita, Jorge
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             CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE
STREET: P.O. Box 4433
                                                                  TITLE OF INVENTION: Compositions and Methods Generation of Transgenic
                                                        NUMBER OF SEQUENCES: 51
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CITY: Houston
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Query Match:
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TELEPAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/949,155
FILING DATE: <Unknown>
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                     ValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGlu 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA
                                        CCCTCCTGGCGAACCGTGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG 2776
                                                                                                                                                                                                                                                                                  TrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrp
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                                                                                                                                                                                                                                                                                                                           GAGCCGGGG-----CCGCCGCCGGAGGTGCACGTGTGGTGGGAGGAGCCCAAG
                                                                                                                                                                                                             GTGCAGTCCCTGTCTGACCAAGTGCAGGAGGAGGTGCTCAGCACCAAGGTCACCCAGGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/819,964 FILING DATE: 28-Mar-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 77210-4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
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Matches:
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Indels:
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RESULT 6
US-08-726-306A-28
                                                                                                       Sequence 28, Application US/08726306A Patent No. 5958684 GENERAL INFORMATION:
                                                    APPLICANT:
APPLICANT:
APPLICANT:
                APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
TITLE OF INVENTION: DIAGNOSIS METHOD AND REAGENTS
NUMBER OF SEQUENCES: 189
   CORRESPONDENCE
                                                                                                                                                                                                                                         ValProSerAspAsn 316
                                                                                                                                                                                                                                                                                                                  GlnTrpAlaGlyLeuValGluLysValGlnAlaAlaValGly---ThrSerAlaAlaPro
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                                                                                        van Leeuwen, Frederik Willem
   ADDRESS
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FILING DATE: 01-Jan-1996
ATTORNEY/AGENT INFORMATION:
NAME: WILLIAMS, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 96,048-A (3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
TELEPHONE: (617) 345-9111
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH. 660 back page 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette, 3.50 inch, COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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STREET: 1 Financial Center
CITY: Boston
STATE: MA
COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                       ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140
                                                                                          ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120
                                                                                                                                                               LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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CGGCTGGGCGGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
                                                                       ACCCCGGTAGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC
                                                                                                                                                CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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                                                       US-09-827-854-17 (1-317) x US-07-849-389-6 (1-330)
                                                                                                              Best Local Similarity:
                                                                                                                           Percent Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6, Application US/07849389
Patent No. 5525493
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 167
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/849,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: HORNES, Erik
APPLICANT: UHLEN, Mathias
TITLE OF INVENTION: CLONING
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 330 base pairs
                                                                                                                                                                                                                                        FEATURE:
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TELEFAX: 899149
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FILING DATE: 19920519
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                             LOCATION:
                                                                                                                                                                                                                        NAME/KEY:
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1800 Diagonal Road,
                                                                                                                                                                                                                                                                                                                                                                                       : (703)836-9300
(703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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US-09-827-854-17 (1-317) x US-08-617-256-24 (1-252)
                                         Query Match:
                                                                                                             Alignment Scores:
                                                                                                                                         US-08-617-256-24
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                                                        Best Local Similarity:
                                                                   Percent Similarity:
                                                                                    Score:
                                                                                                 Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 6043031
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                      TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth A.
REGISTRATION NUMBER: 35,
                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
FILING DATE: March 17, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
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                                                                                                 No.:
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                                                                                                                                                                  STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US, FILING DATE: March 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61
                                                                                                                                                                                                 TYPE:
                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
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60 State Street, suite 510
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                                      5.73e-28
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Conservative:
Mismatches:
Indels:
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US-09-287-141-24
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                 Pred. No.:
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                                                                   US-09-287-141-24
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                                                                                                                                                                          TELECHONE: 619-30 TELEPHONE: 619-587-5360
TELEFAX: 619-587-5360
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
TELEFAX: (617)227-5941
                                                                                                                                                             TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18 MAR-1996
PRIOR APPLICATION DATA:
                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24:
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                              TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                     NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/287,141 FILING DATE:
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                                                                                                    TOPOLOGY:
                                                                                                                                                  LENGTH:
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: CA
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                                                                                                                                                  252 base pairs
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Length:
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Best Local Similarity:
Query Match:
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                                                                                  INFORMATION FOR SEQ ID NO:
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OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:

ADDRESSEE: Heller Ehrman White & McAuliffe
ADDRESSEE: Terming Square, 7th Floor
                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
                                                SEQUENCE CHARACTERISTICS LENGTH: 252 base pairs
                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: K ster, TITLE OF INVENTION:
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                                                                                                                                                                                                 NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002G
                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/6: FILING DATE: 18-MAR-1996
                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62 GTGTGCGC-CGCCTGGTGCAGTACCGCGGCGAGGTGCAGGCCATGCTCGGCCAGAGCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24, Application US/09431613
o. 6221601
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                                nucleic acid
                                                                                                (617)227-7400
(617)227-5941
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Indels:
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US-09-504-245-24
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; MOLECULE TYPE: US-09-431-613-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24, Application US/09504245 Patent No. 6221605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
          TELEFAX: 858-587-5360
INFORMATION FOR SEQ ID NO:
                                                                                                     APPLICATION NUMBER: 08/617
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                               SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatit
OPERATING SYSTEM: DOS
                                     REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002J
TELECOMMUNICATION INFORMATION:
TELEPHONE: 858-450-8400
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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                                   TELEPHONE:
                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                    CITY: La Jolla
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                                                                                          Seidman, Stephanie L
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TOPOLOGY: lir
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Percent Similarity:
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                                                   FILING DATA:
CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/617,256
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
APPLICATION OBTA: 03-MAR-1995
PRICING DATE: 03-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 24, Application US/09287682 Patent No. 6235478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
TITLE OF INVENTION: DNA Diagnostics Based on Mass Spectrometry
NUMBER OF SEQUENCES: 33
                                   FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
                                                                                                                                                                                                                                     CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                   OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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US-09-287-679-24
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1. CURRENT APPLICATION DATA:
                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                        APPLICANT: K ster, Hubert APPLICANT: Little, Daniel P. APPLICANT: Braun, Andreas
                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400
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                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                STREET: 4250 EX CITY: La Jolla STATE: CA
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                                                             APPLICATION NUMBER: FILING DATE:
                                                                                                                     COMPUTER: IBM CONT
OPERATING SYSTEM:
                                                                                                                                                                                    COUNTRY: USA
ZIP: 92037-9103
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619-587-5360
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Matches:
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Best Local Similarity:
Query Match:
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APPLICATION NUMBER: 08/406,199

FILING DATE: 03-MAR-1995

ATTORNEEY/AGENT INFORMATION:

NAME: Seidman, Stephanie L

NAME: Seidman, Stephanie L

NAME: NUMBER: 33,779
                                                                                                                                                                                                                                                                              Sequence 24, Application US/09397766 Patent No. 6268144
                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: K ster, hubert
TITLE OF INVENTION: DNA Diagnostics Based on
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617)227-594 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHAN: 619-587-5360
TELEFAX: 619-587-5360
TELECOMMUNICATION INFORMATION:
                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
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                                                                               COUNTRY: USA
7TP: 92037-9103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               169 AspAlaAspAspLeuGlnLysCysLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla 188
SOFTWARE: PatentIn Release IRRENT APPLICATION DATA:
                                                                                                                             STATE:
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(617)227-5941
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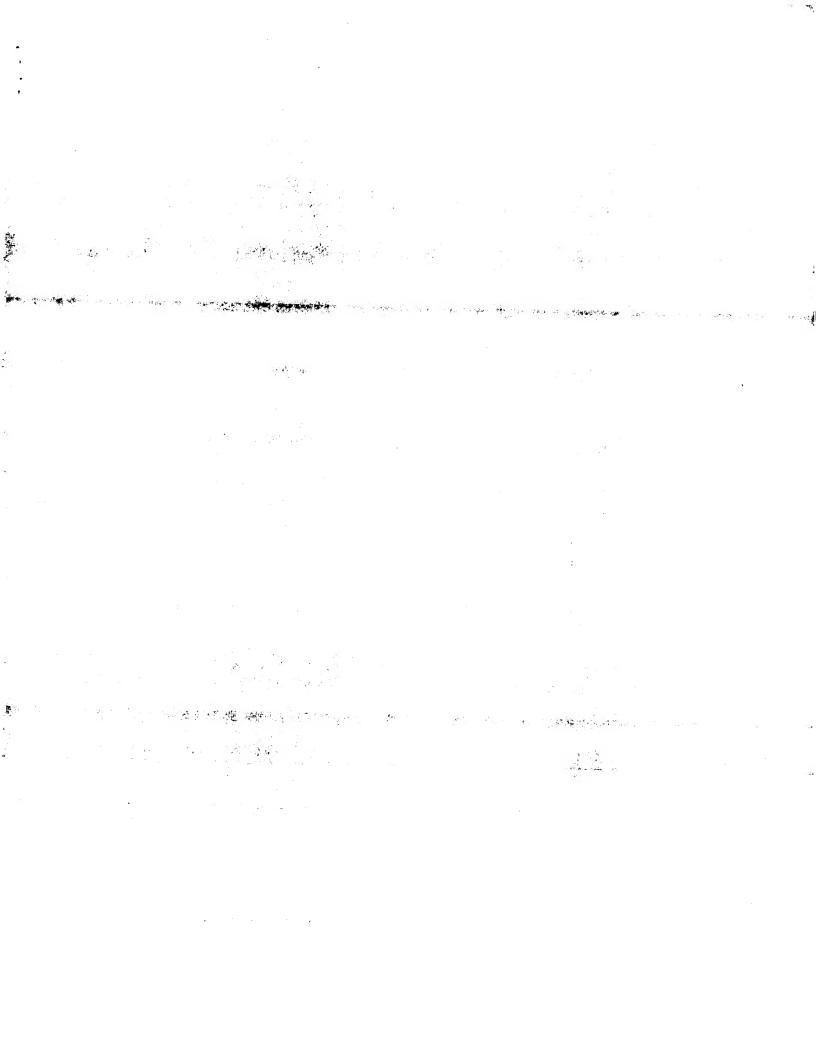
APPLICATION NUMBER:

US/09/397,766

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Best Local Similarity:
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                                                                                                                                      Sequence 24, Application US/09287661
Patent No. 6277573
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                               NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
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TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
                                                                                                              APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
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LENGTH: 252 base pairs
TYPE: nucleic acid
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RECISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002I
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
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APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 08/6:
FILING DATE: 18-MAR-1996
                                                                                                                             APPLICANT:
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CITY: I
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STATE: CA
COUNTRY: USA
ZIP: 92037-9103
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CLASSIFICATION:
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DB:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/(
FILING DATE:
CLASSIFICATION:
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TELEFAX: (617)227-5941
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REGIESTION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Diskette
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TOPOLOGY: 111
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TELEFAX: 619-587-5360
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GluArgGlyLeu 192
                                                                                                      GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGCCCGCGAGGGCCCC
                                                                 AspAlaAspAspLeuGlnLysCysLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla
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SYSTEM: DOS
PatentIn Release #1.0, Version
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Conservative:
Mismatches:
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Search completed: March 14, 2003, 20:18:46 Job time :  $40.7343 \ \text{secs}$ 



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Result
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Maximum
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-MODEL=frame+ p2n.model -DEV=xlp
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-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Title:
Perfect score:
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Ygapop 10.0,
Fgapop 6.0,
Delop 6.0,
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Match
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Listing first 45 s
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3377.460 Million cell updates/sec
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Sequence 10, Appli
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Sequence 17, Appl
                                                                        Description
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US-09-827-854-12
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US-09-954-76-761
US-09-967-013-5
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US-09-967-013-5
US-09-960-352-5420
US-09-960-352-4237
US-09-960-352-4237
US-09-960-352-14047
US-09-960-352-14047
US-09-960-352-14047
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15, Appl 113, Appl 1240, Appl 134, Appl 134, Appl 1400, Appl 1400,
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## PIGNMENT

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GENERAL INFORMATION:

APPLICANT: Zannis, Vassilis

APPLICANT: Kypreos, Kyriakos E.

APPLICANT: Kypreos, Kyriakos E.

TITLE OF INVENTION: Compounds and methods for lowering

TITLE OF INVENTION: Compounds and methods for lowering hypertriglyceridemia

CURRENT APPLICATION NUMBER: US 09/679,088

PRIOR APPLICATION NUMBER: US 09/679,088

PRIOR APPLICATION NUMBER: US 09/544,386

PRIOR APPLICATION NUMBE
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Application

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; Patent No. US20020123093A1
; GENERAL INFORMATION:
APPLICANT: Zannis, Vassilis
; APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lower!
TITLE OF INVENTION: Conbesterol levels without inc
FILE REFERENCE: 07180/004003
; CURRENT APPLICATION NUMBER: US/09/827,854
; CURRENT FILING DATE: 2001-04-05
; PRIOR APPLICATION NUMBER: US 09/679,088
; PRIOR APPLICATION NUMBER: US 09/544,386
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; ORGANISM: Homo
US-09-827-854-9
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SOFTWARE: FastSEQ for Windows
SEQ ID NO 9
LENGTH: 1156
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TITLE OF INVENTION: COMPOSITIONS AND METHODS
FILE REFERENCE: 870759
CURRENT FILING DATE: 2002-01-14
PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 166
SOFTWARE: Patentin version 3.1
SEQ ID NO 129
LENGTH: 1156
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LOCATION: (61)..(1014)
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                                                                                                 SEQ ID NO 17
LENGTH: 1156
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          FEATURE:

NAME/KEY: CDS

LOCATION: (61)...(1014)

OTHER INFORMATION: Nucleotide

OTHER INFORMATION: (APOE)

5-09-802-640-17
                                                                                                                                                                                                                                                    Sequence 17, Application US/09802640 Publication No. US20030036057A1
                                                                                                                                                                                                      APPLICANT: Braun, Andreas
APPLICANT: Bonsal Aruna
APPLICANT: Kleyn Patrick
                                                                                                                          SOFTWARE: FastSEQ
                                                                            ORGANISM: Homo
                                                                                          TYPE: DNA
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CURRENT APPLICATION NUMBER: US/09/802,640 CURRENT FILING DATE: 2001-03-09 NUMBER OF SEQ ID NOS: 122
                                                                                                                                                                                                          TITLE OF INVENTION: GENES AND POLYMORPHISMS TITLE OF INVENTION: CARDIOVASCULAR DISEASE FILE REFERENCE: 24736-2048
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; TYPE: DNA
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APPLICANT: Kypreos, Kyriakos E.
AIPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for the compound of the compounds of the 
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US-09-827-854-8
; Sequence 8, Application US
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; OTHER INFORMATION: Incyte; NAME/KEY: unsure; LOCATION: 46; OTHER INFORMATION: a, t, c US-10-044-090-454
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NUMBER OF SEQ ID NOS: 850
SOFTWARE: PERL Program
SEQ ID NO 454
                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: GENES DIFFERENTIALLY EXPRESSED FILE REFERENCE: PA-0028 US CURRENT APPLICATION NUMBER: US/10/044,090
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                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
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                 SEQ ID NO 12
LENGTH: 1156
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ORGANISM:
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
FILE REFERENCE: (Cholesterol levels without inducing
FILE REFERENCE: (7180/004003

CURRENT APPLICATION NUMBER: US/09/827,854

CURRENT FILING DATE: 2001-04-05

PRIOR APPLICATION NUMBER: US 09/879,088

PRIOR APPLICATION NUMBER: US 09/579,088

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-04-06

PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                                                                  Sequence 12, Application U Patent No. US20020123093A1
                                                                                                                                                                                                                                                APPLICANT: Zannis, Vassilis APPLICANT: Kypreos, Kyriak
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 1156
TYPE: DNA
GRGANISM: Homo sapiens
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Query Match:
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Pred. No.: 1.03e-135 Score: 1561.00 Percent Similarity: 98.748 Best Local Similarity: 98.748 Query Match: 98.248  Query Match: 98.248  Length: 1157 Matches: 0  Mosmatches: 0  Mismatches: 0  Mismatches: 0	ORGANISM: Homo sapiens US-09-954-456-760 Alignment Scores:	SOFTWARE: PatentIn ver SEQ ID NO 760 LENGTH: 1157 TYPE: DNA	FILING DATE: 2000-09-27 APPLICATION NUMBER: US/60/235, FILING DATE: 2000-09-27 R OF SEO ID NOS: 2276	PRIOR FILING DATE: 2000-09-27 PRIOR APPLICATION NUMBER: US/60/235, PRIOR FILING DATE: 2000-09-27 PRIOR APPLICATION NUMBER: US/60/235.	PRIOR FILING DATE: 2000-09-26 PRIOR APPLICATION NUMBER: US/60/235 PRIOR FILING DATE: 2000-09-26 PRIOR APPLICATION NUMBER: US/60/235	PRIOR EILING DATE: 2000-09-25 PRIOR APPLICATION NUMBER: US/60/ PRIOR FILING DATE: 2000-09-25 PRIOR APPLICATION NUMBER: US/60/ PRIOR APPLICATION NUMBER: US/60/	LING DATE: 2000-09-18 PLICATION NUMBER: US/60/ LING DATE: 2000-09-20 PLICATION NUMBER: US/60/	REFE NT A	GENERAL INFORMATION: APPLICANT: Young, Paul FITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Cancer Title OF INVENTION: Sets	RESULT 9 US-09-954-456-760 ; Sequence 760, Application US/09954456 : Datent No [RESON 201150574]	Qy 301 ValGlnAlaAlaValGlyThrSerÅlaAlaProValProSerAspAsnHis 317	Qy 281 SerTrpPheGluProLeuValGluÄspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300	Qy 261 LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 280 	Qy 241 SerArgThrArgAspArgLeuAspoluValLysGluGlnValAlaGluValArgAlaLys 260	Qy 221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240 	Qy 201 ProLeuValGluGlnGlyArgVaiArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220	Qy 181 GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200
RESULT 10 US-09-880-107-2244 US-099-80-107-2244; Application US/09880107 ; Sequence 2244, Application US/09880107 ; Patent No. US20020142981A1 ; GENERAL INFORMATION:	Qy 301 ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 317	Qy 281 SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300 	Qy 261 LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 280 	Qy 241 SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys 260	Qy 221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240	Qy 201 ProLeuvalGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220	Qy 181 GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200 	Qy 161 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr 180 	Oy 141 GlnAlaMetLeuAspGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160 	OY 121 ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140	OY 101 ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120	QY 81 LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluLeu 100	Qy 61 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80	OY 41 G1yG1nArgTrpGluLeuAlaLeuG1yArgPheTrpAspTyrLeuArgTrpValG1nThr 60 	Qy 21 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40	Oy 1 MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20 	DB: 10 Gaps: 0 US-09-827-854-17 (1-317) x US-09-954-456-760 (1-1157)

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FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 2244
LENGTH: 1157
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US-09-880-107-2244
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APPLICANT: Scherf, I
APPLICANT: Gene Log
TITLE OF INVENTION:
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Scherf, Uwe
Gene Logic, Inc.
VENTION: Gene Expression Profiles
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
FITTLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Coholesterol levels without inducing
FILE REFERENCE: 07180,004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 1156
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; ORGANISM: Homo
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Sequence 7, Application US/09827854
Patent No. US20020123093A1
GENERAL INFORMATION:
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Alignment Pred. No. Score: Percent S	RESULT US-09- Sequ Fate Pate APP FIT	ОУ	Qу	Оy	Qу	ОУ	Qу	Qy Db	Qу	Qу	Qy Db	Qy Db	Qу
ent Score No.: t Similar	SULT 12 -09-967-013-5 Sequence 5, Ap Patent No. US2 GENERAL INFORM APPLICANT: St TITLE OF INVE FILE REFERENC CURRENT APPLI CURRENT PILIN PRIOR APPLICAD PRIOR FILING PRIOR FILING PRIOR FILING OMBER OF SEQ SOFTWARE: FOR SEQ SOFTWA	301 Val     961 GTG	281 Ser     901 AGC	261 Leu       841 CTG	241 Ser      781 AGC	221 Leu      721 CTA	201 Pro      661 CCC	181 Gln     601 CAG	161 Lys     541 AAG	141 Gln     481 CAG	121 Arg      421 CGG	101 Thr      361 ACC	81 Leu     301 CTC
res: arity:	0.013-5  5. Applic  40. US20020 INFORMATIC INFORMATIC STANT: Stanto DF INVENTIC FIVENTIC FIVENTIC FAPLICATIO FAPLICATIO FAPLICATIO FAPLICATIO FAPLICATIO FAPLICATIO FAPLICATIO FAPLICATIO FAPLICATIO OF SEO ID	GlnAlaA         CAGGCTG	TrpPheG         TGGTTCG	GluGluG         GAGGAGC	ArgThra	GlnGluA          CAGGAGC	LeuValo           CTGGTGG	AlaGlyA          GCCGGGG	LeuArgI         CTGCGTA	AlaMetI          GCCATGC	LeuGly&         CTGGGCG	Provala           CCGGTGC	MetAspo          arggaco
1.11e-114 1361.50 59.80%	RESULT 12 Scquence 5, Application US/09967013 Sequence 5, Application US/09967013 Patent No. US20020045840A1 GENERAL INFORMATION: APPLICANT: Stanton, Jr, Vincent P TITLE OF INVENTION: METHOD FOR GENETIC AI TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: 11926-022001 CURRENT APPLICATION NUMBER: US/09/967,01: CURRENT FILING DATE: 2000-10-25 PRIOR APPLICATION UNMBER: 60/206,613 PRIOR FILING DATE: 2000-05-23 NUMBER OF SEQ ID NOS: 91 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 5 LENGTH: 41907 TYPE: DNA ORGANISM: Homo sapiens US-09-967-013-5	laValGlyThrSerAlaA 	luProLeuValGluAspM 	lnAlaGlnGlnIleArgI                 AGGCCCAGCAGATACGCC	.rgAspArgLeuAspGlu\                   GCGACCGCCTGGACGAGG	.rgAlaGlnAlaTrpGlyc 	luGlnGlyArgValArg <i>t</i> 	laArgGluGlyAlaGlu# 	ysArgLeuLeuArgAsp <i>t</i> 	.euAspGlnSerThrĠluc                  TCGGCCAGAGCACCGAGC	laAspMetGluAspValc	laGluGluThrArgAla <i>P</i>                   CGGAGGAGCGCGGCAC	luThrMetLysGluLeuI               AGACCATGAAGGAGTTG <i>I</i>
Length: 41 Matches: 29 Conservative: 0	TIC ANALYSIS OF APOE OF 67,013 3	ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 	SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 	LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys 	SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys	LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 	roLeuValGluGlnGlyArgValA <sup>L</sup> gAlaAlaThrValGlySerLeuAlaGlyGlnPro 	lnalaGlyalaArgGluGlyalaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 	LysLeuArgLysArgLeuLeuArgÅspAlaAspAspLeuGlnLysCysLeuAlaValTyr 	lnalametLeuAspGlnSerThr¢luGluLeuArgValArgLeuAlaSerHisLeuArg           	ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 	ThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu 
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US-09-9
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CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver: 2.0
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                                                                                                                                                                                                                                  SEQ ID NO 133
LENGTH: 786
                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 133, Application US/09925302
Patent No. US20020044941A1
                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins
FILE REFERENCE: PAIO4
                                                                                                                                                                                                                                                                                                                                                                                                                  -09-925-302-133
                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo
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                                        ArgValArgLeuAlaSerHisLeuArgLysLeuArgLysArgLeuLeuArgAspAlaAsp
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CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR ETLING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR ETLING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR ETLING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEO ID NOS: 583
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 374
LENGTH: 478
              Qy
                                                                                                                                                                                                       NAME/KEY: misc_feature; LOCATION: (1)...(478); OTHER INFORMATION: n-a,t,g
                                              US-09-827-854-17 (1-317) x US-09-964-824A-374 (1-478)
                                                                                            Query Match:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                            No.:
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MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
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Best Local
Query Match
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APPLICANT: Vockley, Joseph G.
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/21,379
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR APPLICATION NUMBER: US 60/237,054
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2491
LENGTH: 478
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                                                                                                                                                                                        09-880-107-2491
                                                                                                                                                                                               OTHER INFORMATION: Genbank Accession NAME/KEY: unsure LOCATION: (1)...(478)
OTHER INFORMATION: n = a or c or g or
                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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Search completed: March 15, 2003, 03:04:37 Job time: 82.8753 secs

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-WARN_TIMEOUT=30 -THREADS=1 -XGAPD=10 -XGAPDEXT=0.5 -FGAPDP=6 -FGAPEXT=7
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Listing first 45 s
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Maximum DB seq length: 200000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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## ALIGNMENTS

AUTHORS TITLE	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	в1670367	RESULT 1
NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1027)	Homo sapiens	human.	EST.	BI670367.1 GI:15584600	mRNA sequence. BI670367	603292738F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5312024 5',	BI670367 1027 bp mRNA linear EST 12-SEP-2001		

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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), if
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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/lab_host="DH10B"
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EST.
                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
Tissue Procurement: ATCC/DCTD/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLCM1718 row: b column: 10
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Contact: Robert Strausberg,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/clone_lib="NIH_MGC_49"
/tissue_type="melanotic melanoma,
/lab_host="DH10B (phage-resistant)
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
Toshiyuki and Piero Carninci (RIKEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 933)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  603295681F1 NIH_MGC_96 Homo sapiens mRNA sequence.
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/lab_host="DH10B"
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/clone_lib="NIH_MGC_96"
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/db_xref="taxon:9606"
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                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                     Unpublished (1999)
Contact: Robert St
                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (1) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information
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CAGGCCGGGGCCCGAGGGCGCGAGCGCGTGCTCAGCGCCATCCGCGAGCGCCTGGGG
               GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysCysLeuAlaValTyr
                                                                                                                                   CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
                                                                                                                                                   GlnAlaMetLeuAspGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                                                                                                                                      ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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http://image.llnl.gov
Plate: LICM1674 row: c colu
High quality sequence stop: 7
Location/Qualifiers
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/lab_host="DH10B (phage-resistant):
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
EcoRI; cDNA made by Oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GCACGAG(G). Size-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-CDNA synthesis kit (Stratagene) and Superscript
II RT (Life Technologies). Note: this is a NIH_MGC
Library. | "
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/db_xref="taxon:9606"
/clone="IMAGE:4841411"
/clone_lib="NIH_MGC_49"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI),
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11762 row: a column: 11
High quality sequence stop: 762.
Location/Qualifiers
1. 942
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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                                                                                      /note="Organ: brain; Vector: pBluescript (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length [clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

284 c 361 g 120 t
                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
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/clone="IMAGE:5301010"
/clone_lib="NIH_MGC_96"
                                                                                                                                                                                                                                        /tissue_type="hypothalamus"
/lab_host="DH10B"
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                                                                                        CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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Tissue Procurement: DCTD/DTP
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
Unpublished (1999)
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Plate: LLCM1875 row: i column: 03
High quality sequence stop: 792.
Location/Qualifiers
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Clone distribution: MGC clone distribution information can be
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Tissue Procurement: ATCC
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Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incytte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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NIH-MCC http://mgc.nci.nih.gov/.
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BG774871.1
                                                                             cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1612 row: e column: 20 High quality sequence stop: 813.
                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; F
Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 817)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
                                                                                                                                                                                   Contact: Robert Strausberg, I
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4760827"
/clone_lib="NIH_MCC_40"
/tissue_type="carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
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CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
                                                                   GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTTGGGATTACCTGCGCTGGGTGCAGACA
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAMI1726 row: g column: 10
High quality sequence stop: 783.
Location/Qualifiers
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/clone_lib="NIH_MGC_96"
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/db_xref="taxon:9606"
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/lab_host="DH10B"
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High quality sequence stop: 829.
Location/Qualifiers
                                                                                                                                                                                                    CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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602763768F1 NIH_MGC_42 Homo sapiens cDNA clone IMAGE:4899112 5',
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National Institutes of Health, Mammalian
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                                                                                                                   1. .845
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Directionally cloned into ECORI/XhoI sites using the
following 5 adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Lir
Hong in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-CDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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SLeuGluGluGln :
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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Plate: LLAM10672 row: e column:
High quality sequence stop: 781.
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Contact: Robert Strausberg,
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National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                            mRNA sequence.
BI551475
BI551475.1 GI
                                                                                                  Email: cgapbs r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI)
Toshiyuki and Piero Carninci (RIEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAMI1691 row: 1 column: 04
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603194314F1 NIH_MGC_95
                                                                                                                                                                                                                       Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                             NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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Location/Qualifiers
1. 919
                                                                                                                                                                                                                                                                                                         sapiens
/organism="Homo sapiens"
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/clone="IMAGE:5274003"
/clone_lib="NIH_MGC_95"
/tissue_type="hippocampus
/lab_host="DH10B"
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TITLE
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GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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1. (bases 1 to 757)
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National Institutes of Health, Mammalian
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http://image.llnl.gov
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/db_xref="taxmon:0006"
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/clone="tMAGE:5557004"
/clone="the-"NIH_MGC_112"
/tissue_type="melanotic melanoma, cell line"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mRNA sequence.
BG706129
BG706129.1 GI:
EST.
                                                                                                                                                                                                                   Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                   High
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602669093F1 NIH_MGC_96 Homo sapiens
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National Institutes of Health, Mammalian Gene Collection (MGC)
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http://image.llnl.gov
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/clone_lib="NIH_MGC_96"
                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
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Command line parameters:

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-Q-/cgn2_1/USPT0_Spool_VG09827854/runat_11032003_101610_27486/app_query.fasta_1.3576

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-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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                                                                                Apó E humanized mammal
Patent: JP 2001017028-A 2 2:
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/2
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128:
PR SHINOBU FUJITA, HIROKI I
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                                                         PI SHINOBU FUJITA,HIROKI HAMANAKA,YUKO FUKUI,MINESUKE AOLIKG7/027,A61K45/00,A61P25/28,A61P43/00,C12N5/10, PC C12N15/09/C07K14/775, PC (C12N5/10,C12R1:91),C12N5/00,C12N15/00,(C12N5/00,C1
                                                                                                                                                           Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 1156)
Pujita,S., Hamanaka,H., Fukui
                                                                                                                                                                                                              Apo E humanized mammal.
BD004278
BD004278.1 GI:18632239
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               Location/Qualifiers (61) (1011).
Location/Qualifiers
1. 1156
/organism="Homo sapiens"
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Isolation and Characterisation of a variant allele of the human apoliopprotein E Biochem. Biophys. Res. Commun. 130 /3:
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84185684
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Type III hyperlipoproteinemia associated with apolipoprotein phenotype E3/3. Structure and genetics of an apolipoprotein
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The isolation of cDNA clones for human apolipoprotein E and the detection of apoE RNA in hepatic and extra-hepatic tissues EMBO J. 2 (12), 2369-2373 (1983)
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83082756
[6] sites; mutations resulting in type III hyperlipoproteinemia. Apo E is a component of normal human very low density lipoprotein. There are six human apo E phenotypes known to result from a single structural gene, three of the common alleles being epsilon-4,
                                                                epsilon-3 and variant. [5] epsilon-2 allele.
                                                                                                                              J. Clin. Invest.
89198059
2539388
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Taylor, J.M.
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Zannis, V.I., McPhers
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. Chem. 258, 11422-11422 (1983)]
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Apo E is located on chromosome 19 --Jackson, Bruns and Breslow , PNAS USA 81, 2945-2949 (1984) -- and is believed to be linked to the Apo C-II gene (see separate entry).

The two mutations causing type III hyperlipoproteinemia (apolipoprotein E phenotype E3/3) produces substitutions of Arg for Cys at amino acid 112 and Cys for Arg at amino acid position 142. Draft entry and printed copy of sequence for [1] kindly provided by L.L. Gill, University of Aberdeen.

Complete source information:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          see the separate entry. [J. Biol. Chem. 258, 11422-11422 (1983)] is too extensively revised relative to [1] to record the revisions in the FEATURES table. The sequence below is that of [J. Biol. Chem. 258, 11422-11422 (1983)]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      epsilon-3 and epsilon-2. This sequence appears to be of the epsilon-3 allele. [1] argues that the apo E polymorphism involves mutations in the structural coding region; for example the epsilon-2 phenotype which is characterized by hyperlipoproteinemia is thought to result from a c to t change (arg to cys) at base 586 below [3], [5]. The sequence shown is 57% homologous with human apo
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/replace="a"
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                                                               GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                                        GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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          cDNA Library Preparation: Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Con DNA Sequencing by: Genome Sequence Centre, BC. Cancer Agency, Vancouver, BC, Canada
                                                                                                                                                                    Submitted (20-FEB-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                             Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1186)
                                                                                 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                   NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens, complete cds.
info@bcgsc.bc.ca
                                                                                                                                                                                                                           Direct Submission
                                                                                                                                                                                                                                           Strausberg,R.
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MGC:1571 IMAGE:3355712, mRNA,
                                                   Consortium
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Jones,

Jennifer

Asano,

Ian Bosdet,

Yaron

Butterfield

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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                                                                                                                                                                                                                   LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                                                                                                                                                                                                                                                                             ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
                                                                                                                                                                                                                                                                                                                     GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
ArgLeuGlyAlaAspMetGluAspValÇysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                           CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
                                                                                                                                                                                                                                                                                                  GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                              ACCCCGGTGGCGGAGGAGACGCGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
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This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 4557324.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy, George Yang, Scott Zuyderduyn, Marco Marra.
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protein_id="AAH03557.1"

protein_id="AAH03557.1"

/protein_id="AAH03557.1"

/db_xref="fd:13907699"

/translation="MKVLWAALLVTFLAGCQAKVEQAVETEPEPELRQOTEWQSGQRW

ELALGREWDYLRRVQTLSEQVOELLSSQVTOBELRALMDETMELKAYKSELLEBOLTP

VAEETRARLSKELQAAQARLGADMEDVGGRLVQYRGBVQAMLGQSTEBLKAVRULASHLF

VAEETRARLSKELQAAQARLGADAREGAERGLSAIRERLGPLVEQGRVVAATVGSLAG

OPLQERAQAWGEELRARNEEMGSRTROBILDEVKEQVAFQVRAKLEEQAQQIRLQAEAFQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="Bye, retinoblastoma"
/clone_lib="NIH_MGC_16"
/lab_host="DH10B-R"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="LocusID:348"
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/clone="MGC:1571 IMAGE:3355712"
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                                  Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

Elikaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

Eli (bases 1 to 1110)

Elikaryota; Metazoa; Chordata; Craniata; Vertebrata; Homo.

Elikaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

Elikaryota; Metazoa; Catarrhini; Hominidae; Homo.

Elikaryota; Metazoa; Catarrhini; Hominidae; Homo.

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Elikaryota; Metazoa; Catarrhini; Hominidae; Homo.

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JP 1994315392-A/1.
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                        Location/Qualifiers
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/db_xref="taxon:9606"
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Mammalia; Eutheria; Primates; C:
1 (bases 1 to 1156)
Fujita,S., Hamanaka,H., Fukui,Y
Apo E humanized mammal
Patent: JP 2001017028-A 1 23-JAI
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/1
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128919
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Apo E hur
BD004277
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JP 2001017028-A/1.
Homo sapiens.
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Location/Qualifiers 1. .1156
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/db_xref="taxon:9606"
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          Similarity:
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Sequence 378
AX333278
AX333278.1
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Eukaryota; Me
Mammalia; Eut
                                                                                                                                    Cancer gene determination
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                                                                     212
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Pharmaceuticals (US
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                                                                                        Location/Qualifiers
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                                                                    ; Metazoa;
Eutheria;
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99.05%
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Patent WO0194629.
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                                                                            sapiens"
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Length:
Matches:
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                                                           ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
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Gene expression profiles in liver cancer
Patent: WO 0229103-A 2244 11-APR-2002;
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                                                                                                                                                                                                                                                      to mRNA.
                                                                                                                                                                                                                                                                sapiens (clone: pHAE[112,178,813].) male 57-year
                                                                                                                                                      Chem.
 /gene="APOE"
<1. .1157
                    /tissue_type="liver"
/dev_stage="57-year old"
1. .1157
                                                     /clone="pHAE[112,178,813]."
/sex="male"
                                                                           /map="19q13.2"
                                                                                      /organism="Homo|sapiens"
/db_xref="taxon:9606"
                                                                                                                      Location/Qualifiers
                                                                                                                                                      259 (10),
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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                                                                                                                   GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                                                                     GAGCAAGCGGTGGAGACAGAGCCGGAGCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
                                                                                                                                                                                      GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                  CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
                                                                                                   GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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/protein_id="AAB59518.1"
/protein_id="AAB59518.1"
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QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH"
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/replace="g"
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/replace="a"
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/note="a in pHAE[112,178];
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JP 2001017028-A/3.
Homo sapiens
Homo sapiens
            PT SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE YOKOYAMA
A01K67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC
C12N15/09//C07K14/775,
PC (C12N5/10, C12R1:91), C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
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BD004279
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1 (bases 1 to 1156)
Fujita,S., Hamanaka,H., Fukui,Y. and Yokoyama,M.
                                                                                                                                                                                                                                                                                                                            humanized
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                                                                                                                                                                                                                                                                                               LysLeuCysLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
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RESULT 15 AF261279 AF261279 AF261279 AF261279 DEFINITION AF261279 VERSION AF261279 VERSION AF261279 VERSION AF261279 VERSION AF261279.1 GI:11034800 KEYWORDS SOURCE Homo sapiens ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Qy 281 SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys 300		101 6 623 C 201 F 683 C	141 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg	Oy 61 LeuSerGluGlnValGlnGluGluEuLeuSerSerGlnValThrGlnGluLeuArgAla 80	QY 21 GLUCHMAL CYALIS LUTING THE FOUND CONTROL OF THE POLICY OF THE POLI	

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 5491)

Nickerson,D.A., Taylor,S.L., Fullerton,S.M., Weiss,K.M.,
Clark,A.G., Stengard,J.H., Salomaa,V., Boerwinkle,E. and Sing,C.F.
Sequence diversity and large-scale typing of SNPs in the human
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Nickerson, D.A.
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Genome Res. 10 (10), 1532-1545 (2000)
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-USER-US09827854 -GCGN_1_1_1201_@frunat_11032003_101609_27476 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NUG_ISCORES=0 -WAIT -LONCLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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hypertriglyceridaemia; lo
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05-APR-2001;
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                                                                                                                                                                                                                                               The patent discloses novel genetic variants of human apolipoprotein CC E (APOE) gene. The invention also relates to compositions and methods CC for haplotyping and/or genotyping the APOE gene. The haplotyping CC methods of the invention are useful for improving the efficacy and CC reliability of several steps in the discovery and development of CC drugs for treating diseases associated with APOE activity, e.g. CC familial dysbetalipoproteinaemia, type III hyperlipoproteinaemia, CC atherosclerosis, and Alzheimer's disease. They are useful to validate CC APOE as a candidate agent for treating a specific condition or disease predicted to be associated with APOE activity and in the design of C clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with APOE activity. Genotyping CC or haplotyping methods are useful to screen for compounds targeting APOE to treat a specific condition or disease associated with APOE activity. The present sequence is a cDNA encoding human APOE protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene -
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GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                          GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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                                                                                                                      Detecting and identifying ovarian tumor, identifying increased risk developing ovarian cancer, and determining effectiveness of ovarian cancer treatment, by measuring expression level of ovarian tumor mar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumours in an individual via the detection and measurement of the expression of ovarian tumour marker genes (ABA83081-ABA83122, ABA83180, ABA83182 and ABA83184) or segments thereof (ABA83123-ABA83169, ABA83179, ABA83181 and ABA83183). The methods of the invention are useful for detecting an ovarian tumour in a patient, for identifying an individual at increased risk for developing ovarian cancer, in prognostic tests for assessing the relative severity of ovarian cancer, in tests for monitoring a patient in remission from ovarian cancer and in tests for monitoring disease status in a patient being treated for ovarian cancer. The methods can additionally be used to identify a particular tumour as being an ovarian tumour (i.e., an epithelial ovarian tumour selected from serous cystadenoma, borderline serous tumour, serous cystadenocarcinoma, mucinous tumour tumour control to the control of the co
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The present invention relates to an ApoE humanised mammalian cell. The present sequence is the coding sequence for human ApoE3, which was used in the method of the present invention. The ApoE humanised mammalian cecan be used for screening for agents useful for treating or preventing
                                                                                                                    Disclosure;
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/product= "Human ApoE3"
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04-OCT-2000;
05-APR-2001;
                                                                                                                                                                                                            The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LI receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia
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P-PSDB; AAE13294.
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hypertriglyceridaemia; low density lipoprotein;
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                                               SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                                                                       GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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                                                                                                                                   SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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                                                      Best Local Similarity:
Query Match:
DB:
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                                                                                                                                                The present sequence is a human apolipoprotein E (apoE) allele, apoE2** DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LI receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
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04-OCT-2000;
05-APR-2001;
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                                                                                                                                  Sequence
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(UYBO-)
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hypertriglyceridaemia; lc
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Location/Qualifiers 61..1014 /*tag= a
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                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to an ApoE humanised mammalian cell. The present sequence is the coding sequence for human ApoE2, which was use in the method of the present invention. The ApoE humanised mammalian can be used for screening for agents useful for treating or preventin Alzheimer's disease and arteriosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
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New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia Claim 14; Page 81-82; 91pp; English.

The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE2 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LDL) receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or having a defect in remnant removal.

Sequence 1156 BP; 208 A; 367 C; 432 G; 149 T; 0 other;

Percent Similarity:
Best Local Similarity:
Query Match:
DB: Scores: 1.31e-114 1569.00 99.37% 99.37% 99.37% 98.74% Conservative: Mismatches: Indels: Gaps: Matches: 1156 315 0 2 0

US-09-827-854-18 (1-317) x AAD22049 (1-1156)

240	$\tt LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly: \\$	221	Qy
720		661	DЪ
220	ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro	201	QΥ
660		. 601	DЪ
200	GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly	181	Qy
600		541	Db
180	LysLeuCysLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr	161	Qy
540		481	Дb
160	GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg	141	Qy
480		421	Db
140	ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal	121	Qy
420		361	рb
120	ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla	101	Qy
360		301	рb
100	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu	81	у
300	CTGTCTGAGCAGGTGCAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG	241	DЬ
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60	GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr	41	Qy
180		121	Db
40	GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer	21	QΥ
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       Recombinant human apolipoprotein-E (ApoE) (AAR86791) can be produced by preparing a genetic construct (contg. an ApoE-encoding sequence, e.g. AAT06957, and flanking regulatory sequences enabling the protein to be expressed in insect cells), which is then introduced into a Manduca sexta larva (using a recombinant Autographica californica nuclear polyhedrosis virus) and recovering the protein from the haemolymph of the larval host. The ApoE produced is in a form sufficiently complexed with lipids to be biologically active, which cannot be achieved in insect cell cultures, and can therefore be
                                                                                                        Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel method for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma or metastatic liver tumour in a patient, and differentiating metastatic liver cancer from the patocellular carcinoma in a patient, involving detecting the level of expression of two or more genes represented in ABN93503-ABN97455 in a ctissue sample. The method of the invention has hepatotropic, and cytostatic activity. The method is useful for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma and metastatic liver carcinoma in a patient. The method is useful for identifying expression profiles which serve as useful diagnostic markers as well as markers that can be used to mobitor disease states, disease progression, conducted the progression, but was obtained in electronic form part of the printed specification, but was obtained in electronic format directly from WIPO at the proposition of the proposition of the printed specification, but was obtained in electronic format directly from WIPO at the proposition of the printed proposition in the proposition of the printed specification, but was obtained in electronic format directly from WIPO at the proposition of the printed proposition in the proposition in the proposition of the printed proposition in the proposition in the proposition of the printed proposition in the proposition in the proposition of the printed proposition in the proposition of the printed proposition in the proposition in the proposition of the printed proposition in the proposition in the proposition of the printed proposition in the proposition in the printed pr
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                                                                                                                                                                                                                                                                                                                           CC progression of benign prostatic hyperplasia (BPH), or screening (II) for CC or identifying an agent that modulates the onset or progression of BPH. CC The method is based on changes in gene expression in BPH tissue isolated CC from patients exhibiting different clinical states of prostate CC hyperplasia as compared to normal prostate tissue. (I) comprises CC detecting the expression levels of one or more genes in prostate cells (II) comprises preparing a first gene expression profile CC prostate cells. (II) comprises preparing a first gene expression profile CC agent, preparing a second gene expression profile of the equal to the CC agent, preparing a second gene expression profile of EPH cells or BPH-like cell population, exposing the cells to the CC cells, and comparing the first and second gene expression profiles. (I) is useful for diagnosing the onset or progression of BPH. (II) is ceful for diagnosing the onset or progression cC the expression level in a tissue or cells, by comparing the expression CC the expression of gene in the database, and displaying the compared to the expression level in BPH. Agents using (II) are useful for treating BPH or prostate cancer. ABK64106-ABK64860 represent humann CC benign prostatic hyperplasia gene sequences of the invention.
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30-MAY-2001; 2001WO-US10838

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The present invention describes a method (M1) for screening for an anti-neoplastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change in capression of at least one gene (I) of a signature gene set, where (I) comprises a sequence (S) selected from 8447 sequences (given in ABL61664 to ABL70110), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has cytostatic activity and can be used in gene therapy. M1 can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a cresult of M1, and the data is sufficient to convey the chemical treatment of cancer such as colon, breast, stomach, lung, thyroid, coesophageal, ovarian, kidney, prostate or pancreatic cancer, infiltrating lobular cancer, squamous cell cancer, infiltrating ductal cancer, carcinoma, papillary carcinoma and Wilm's tumour.
                                                                                                                                                                                                                                                                                                                                                                              Screening for anti-neoplastic agent involves exposing cells to chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature generation.
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GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                     ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
                                                                                                                                                             LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
                                                                                                                                                                                                                                                                                           GTGCAGGCTGCCGTGGGCACCAGCGCCCCCCTGTGCCCCAGCGACAATCAC
                                                                                                                            SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
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uAlaGlyGlnPro 220 	AlaAlaThrValGlySer	ProLeuValGluGlnGlyArgVal	Qy 201	
gGluArgLeuGly	1GluArgGlyLeuSerAlaIleAr 	GlnAlaGlyAlaArgGluGlyAla	Qy 181 Db 601	
gLeuAlaValTyr              CCTGGCAGTGTAC	'gAspAlaAspAspLeuGlnLysArg 	LysLeuCysLysArgLeuLeuAr	Qy 161 Db 541	
aSerHisLeuArg 1              CTCCCACCTGCGC 5	31uG1uLeuArgVa1ArgLeuA1a: 	GlnAlaMetLeuGlyGlnSerThrG 	Qy 141 Db 481	
rArgGlyGluVal 1              CCGCGGCGAGGTG 4	ValcysGlyArgLeuValGlnTyr 	ArgLeuGlyAlaAspMetGluAspVa 	Qy 121 Db 421	
nAlaAlaGlnAla 12             GGCGGCGCAGGCC 42	AlaĀrgLeuSerLysGluLeuGlnAlaAl 	ThrProValAlaGluGluThrArgA 	Oy 101	
GluLeuGluGluGlnLeu 100 	GluLeulysalaTyrlysSerGlule           agttgaaggcctacaaatcggaact	LeuMetAspGluThrMetLys	Qy 81 Db 301	
rGlnGluLeuArgAla 80                  CAGGAACTGAGGGCG 300	LeuSerSerGlnValTh 	LeuSerGluGlnValGlnGluGluLeu 	Qy 61 Db 241	
gTrpValGlnThr 60                 CTGGGTGCAGACA 240	31 yArgPheTrpAspTyrLeuArg              	GlyGlnArgTrpGluLeuAlaLeuG 	Qy 41 Db 181	
rGluTrpGlnSer 40                CGAGTGGCAGAGC 180	luProGluLeuArgGln 	GluGlnAlaValGluThrGluProg: 	Qy 21 Db 121	
SGlnAlaLysVal 20                CCAGGCCAAGGTG 120	aLeuLeuValThrPheLeuAlaGlyCysG] 	MetLysValLeuTrpAlaAl 	Qy 1 Db 61	
	6 (1-1156)	-854-18 (1-317) x AAF8431	US-09-827	
6	Length: 115 Matches: 315 Conservative: 0 Mismatches: 2 Indels: 0 Gaps: 0	: Scores: 2.7e-114 1565.00 1565.00 1565.00 1565.00 1565.00 199.378 1 Similarity: 99.378 1 Similarity: 98.498 1 Similarity: 22	Alignment Pred. No.: Score: Percent Si Percent Local Query Matc DB:	
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d mammalian cell. The ApoE4, which was use numanised mammalian creating or preventing	to an ApoE humanised sequence for human nvention. The ApoE hagents useful for trosclerosis.	present invention relate ent sequence is the coah he method of the present be used for screening for eimer's disease and arts	CC The CC pres CC in t CC can CC Alzh	
	Japanese.	closure; Page 16-17; 22pp;	PS Disc	
ng for agents e and	cell useful for screening ting Alzheimer's disease	apoE humanized mammalian of treating or preventeriosclerosis -	new use art	
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		ru ) MITSUBISHI CHEM CORP.	PA (MITU	
		MAY-1999; 99JP-0125647.	- 90	

Search completed: March 14, 2003, 12:43:20 Job time : 183.728 secs

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e 3, Appl	Sequence 3, Appli	e 2, Appl	e 4, App	e 1, Appl	e 2, Appl	e 3, Appl	e 3, Appl	e 34,	e 34, App	e 3, Appl	1, Appl	<u>,                                     </u>	e 1, Appl	e 12, App	12, App	e 2,	e 2, Appl	e 5,	14, App	e 3, Appl	2, Appl	7, Appl	e 5, Appl	<ol><li>Appl</li></ol>	e 4,	e 4, Appl	4, Appl	e 1, Appl	e 24,	e 24, App	equence 24, App	4, App

## ALIGNMENTS

RESULT 1
US-07-709-949-1
Sequence 1, Application US/07709949
Sequence 1, Application US/07709949

Patent No. 5472858
GENERAL INFORMATION:
APPLICANT: Attie, Alan D
APPLICANT: Sturley, Stephen L
APPLICANT: Beckage, Nancy E
TITLE OF INVENTION: Production of Recombinant Proteins in TITLE OF INVENTION: Production of Recombinant Proteins in TITLE OF INVENTION: Insect Larvae
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Quarles & Brady
STATE: WI
COUNTRY: USA
ZIP: 53701-2113
COMPUTER: EBM PC COMPATIBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/709,949
FILING DATE: 1991664
CLASSIFICATION UNDER: US/07/709,949
FILING DATE: US/07/709,949
FILI

SEQUENCE CHARACTERISTICS: LENGTH: 1157 base pairs

В Ωy Ъ δÃ 밁 Qy B ρy Percent Similarity:
Best Local Similarity:
Query Match:
DB: COATION: 62..119
PUBLICATION INFORMATION:
CHARTICAL TOWNS MCLean, J W
JOURNAL: Journal of Biological Chemistry
VOLUME: 259
PAGES: 6498-6504
DATE: 1994
RELEVANT RESIDUES IN SEQ ID NO: 1: FROM 1
US-07-709-949-1 Qy δÃ γQ Ş В В Q. DЬ В В US-09-827-854-18 (1-317) x US-07-709-949-1 (1-1157) Alignment Scores: Pred. No.: Score: NAME/KEY:
LOCATION:
FEATURE: MOLECULE TYPE: CI HYPOTHETICAL: NO ANTI-SENSE: NO FEATURE: FRAGMENT TYPE:
ORIGINAL SOURCE: 422 242 181 542 482 362 161 141 121 101 302 182 122 TYPE: NUCLEIC ACID
STRANDEDNESS: double
TOPOLOGY: linear 81 61 41 62 21 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer ORGANISM: NAME/KEY: misc\_signal LOCATION: 62..119 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly CGGCTGGGCGCGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu ACCCCGGTAGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG GAGCAAGCGGTGGAGACAGAGCCCGGAGCCGAGCTGCGCAGCAGACCGAGTGGCAGAGC CDS 62..1015 Homo sapiens N-terminal CDNA to mRNA 1.23e-145 1568.00 99.05% 99.05% 99.05% 98.68% Length:
Matches:
Conservative:
Mismatches:
Indels: Н TO 1157 314 0 0 3 0 601 180 541 160 481 140 421 120 361 100 301 80 241 60 181 40 121 20

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; LOCATION:
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 CTCCTCAAAGGCTGGTTCGAGCCTCTGGTGGAAGACATACGGCGCCAGTGGGCCGGGCTG
                                          CGCACCAAAGTGGAGGAGCAGGGCAGCCAGGTTGCGCCCTGCAGGCCGAGGGGATTCCACGCC
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          ArgLeuLysSerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeu
                                                      GAGATGGGCAGCCGGACCCGCGACCGCCTGGATGAGATGCCGTGAGCAGCTGGAGGAGGTG
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                                                                                                                                 US-09-827-854-18 (1-317) x US-09-819-964-5 (1-1126)
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US-09-819-964-5
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                                                                                                                                                                                                                                                    No.:
111
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FILING DATE: «Unknown»
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK:177
                               21
                                                                  51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE AND DURKEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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Bazer, Fuller W
                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Compositions and Methods
                   GluGlnAlaValGluThrGluProGluProGluLeuArg------GlnGlnThrGlu
                                                                              MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
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                                                                ATGAGGGTTCTGTGGGTTGCTTTGGTGGTAACCCTCCTCGCAGGATGCCGGACAGAGGAC
                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/819,964 FILING DATE: 28-Mar-2001
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Conservative:
Mismatches:
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                                                                                                      Sequence 51, Applicat Patent No. 6271436 GENERAL INFORMATION:
                                                  NUMBER OF SEQUENCES: 5
                                                                                     APPLICANT: Piedrahita, Jore APPLICANT: Bazer, Fuller W
                                                                    TITLE OF INVENTION: Compositions and Methods TITLE OF INVENTION: Generation of Transgenic
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COUNTRY: US
ZIP: 77210-4433
                STATE: TX
                          CITY:
                                  ADDRESSEE: ARNOLD, WHITE AND STREET: P.O. Box 4433
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                         Houston
                                                                                                                        Application US/08949155
                                                                                              Piedrahita, Jorge A
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                                           DURKEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TA
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US
FILING DATE: Concurren
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                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                               2490
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               No.:
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                                                                                                                                                                                                                                                                                                                                                                                                        21 GluGlnAlaValGluThrGluProGluProGluLeuArg------GlnGlnThrGlu
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                       TTGGGGGGAAAAAACTTTTTCTTTTTTTTCTTTCATTTGACTTCATGTCTTGCTTTCCTTTCC
                                                                                     CCCTCCTGGCGAACCGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
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Matches:
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Mismatches:
Indels:
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RESULT 5
US-09-819-964-51
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GENERAL INFORMATION:
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                                                                                APPLICANT: Piedrahita, Jorge A
Bazer, Fuller W
TITLE OF INVENTION: Compositions at
Generation of of the composition of the comp
                       CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE
STREET: P.O. Box 4433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTGAGTACCAGGGCCGGCCAGCCGCTGCGCGAGCGCGGAAGCCTGGGGCCAGAAGCTG
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CITY: Houston
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TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                CCCTCCTGGCGAACCGTGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
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FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: <Unknown>
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994.00
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48.76%
62.56%
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Conservative:
Mismatches:
Indels:
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Sequence 28, Application US/08726306A
Patent No. 5958684
GENERAL INFORMATION:
APPLICANT: Van Leeuwen, Frederik Willem
APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
TITLE OF INVENTION: DIACMOSIS METHOD AND REAGENTS
NUMBER OF SEQUENCES: 189
                                                                                                                                                                                                        ValProSerAspAsn
                                                                                                                                                                                                                                                                                                            GluAlaPheGlnAlaArgLeuLysSerTrpPheGluProLeuValGluAspMetGlnArg
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ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 96,048-A (32.
TELECHONUS: (617) 345-9110
TELEPHONE: (617) 345-9111
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 660 base pairs
TYPE: nucleic acid
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Best Local Similarity:
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MOLECULE TYPE:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
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422
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STREET: 1 Financial Center
CITY: Boston
STATE: MA
COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                          62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/726,306A FILING DATE: 02-Oct-1996
                                                                                                                                                                                                                                                                                              ArgLeuGlyAlaAspMetGluAspValCySGlyArgLeuValGlnTyrArgGlyGluVal 140
                                                                                                                  LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGGCCTGGTGCAGTACCGCGGCGAGGTG
                                                                         ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120
                                                                                                                                                                                                                                       GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                                                                                                                                                                                                                   GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                           ACCCCGGTAGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC
                                                                                                                                                                             CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
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Matches:
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RESULT 6 US-08-726-306A-28

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                                                          US-09-827-854-18 (1-317) x US-07-849-389-6 (1-330)
                                                                                                                                                                                        Alignment Scores:
                                                                                                                                                                                                                       US-07-849-389-6
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                                                                                                                          Best Local Similarity:
                                                                                                                                         Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6, Apprix
Sequence 6, Apprix
Sequence 6, Apprix
Sequence 6, Apprix
                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 1678
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-3900
TELEFAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/07/849,389
FILING DATE: 19920519
CLASSIFICATION: 435
CCASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 330 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: HORNES, Erik APPLICANT: UHLEN, Mathias
                                                                                                                                                                      No . :
                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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TELEFAX: 899149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              161 LysLeuCysLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr 180
                                                                                                                                                                                                                                    NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1800 Dia
CITY: Alexandria
STATE: Virginia
             87 LysGluLeuLysAlaTyrLysSerGluLeuGluGluGluGlnLeuThrProValAlaGluGlu
                                                                                                                                                                                                                                                                                  ORGANISM:
                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
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AAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTGACCCCGGTGGCGGAGGAG
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                                                                                                      Matches:
Conservative:
Mismatches:
Indels:
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                   Percent Similarity:
Best Local Similarity:
                                          Score:
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          Query Match:
                                                                 Alignment Scores:
                                                                                      US-08-617-256-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6043031
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 24,
                                                                                                                                                               TELEFAX: (617)227-594
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                        REFERENCE/DOCKET NUMBER: SQ
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAN: (617)227-5941
                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/40
FILING DATE: MARICH 17, 15
ATTORNEY/AGENT INFORMATION:
NAME: AFFOLIA, Beth A.
                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Reli
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: DNA Diagnost NUMBER OF SEQUENCES: 33 CORRESPONDENCE ADDRESS: ADDRESSE: LAHIVE & COCKFIELD
                                                                                                 MOLECULE TYPE:
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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                                                      No.:
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                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/617,256 FILING DATE: March 18, 1996
                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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CITY: Boston
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                                                                                                           STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                  TYPE:
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                                                                                                                                            LENGTH:
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                                                                                                                               nucleic acid
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                                                                                                                                           252 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                               Length:
Matches:
Conservative:
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US-09-827-854-18 (1-317) x US-08-617-256-24 (1-252)

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RESULT 9
US-09-287-141-24
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                 Pred.
                              Alignment Scores:
                                                               US-09-287-141-24
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                                                                                                                                                            TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                           TELEPHONE: 619-450-8400
TELEPAX: 619-587-5360
TELECOMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/406,199
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002D
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/6: FILING DATE: 18-MAR-1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor
                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                 NO::
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                                                                                                TOPOLOGY: lin
                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/287,141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                  252 base pairs
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                                                                                                 linear
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Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mass
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RESULT 10
US-09-431-613-24
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                    TELEFAX: (617)227-594 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                     NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 247
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                       SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                                                                                    FILING DATE: 03-MAR-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                               PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: K ster, TITLE OF INVENTION:
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             STRANDEDNESS:
                                                                                                 TELEPHONE:
                                                                                                                              TELEPHONE: 619-450-8400 TELEFAX: 619-587-5360
                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 18-MA
                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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STATE: CA
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                        nucleic acid
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                                                                                  : (617)227-7400
(617)227-5941
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DNA Diagnostics Based
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Indels:
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RESULT 11
US-09-504-245-24
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24, Application US/09504245 Patent No. 6221605
                                                                                                                                                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/617,256

FILING DATE: 18-MAR-1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/406,199

FILING DATE: 03-MAR-1995

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                               TELEFAX: 858-587-5360
INFORMATION FOR SEQ ID NO:
                                                                NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002J
TELECOMMUNICATION INFORMATION:
TELEPHONE: 858-450-8400
                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
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STATE: CA
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                                                                                                                                                                                                                                                                                 CLASSIFICATION:
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|VENTION: DNA Diagnostics Based
|EQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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Matches:
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Indels:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24, Application US/09287682 Patent No. 6235478
         APPLICATION NUMBER: 08/406,19
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 2473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                    CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMEER: 08/617,256
FILING DATE: 18-MAR-1996
                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
                                                                                                                                                                                                                                              CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: DNA Diagnostics Based on Mass
TELECOMMUNICATION INFORMATION:
                                                                                                                     PRIOR APPLICATION DATA:
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                                                                                                                                                                                                         APPLICATION NUMBER: US/09/287,682 FILING DATE:
                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 92037-9103
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US-09-827-854-18 (1-317) x US-09-504-245-24 (1-252)
109 AlaArgLeuSerLysGluLeuGlnAlaAlaGlnAlaArgLeuGlyAlaAspMetGluAsp
                                           Mismatches:
Indels:
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Patent No. 6258538
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERTSTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/6:
FILING DATE: 18-MAR-1996
                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1 CURRENT APPLICATION DATA:
                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                             APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                               STATE: CA
COUNTRY: USA
ZIP: 92037-9103
                                                                                                                                                                                                                      STREET: 4250 EXCITY: La Jolla
                                              CLASSIFICATION:
                                                            FILING DATE:
                                                                         APPLICATION NUMBER:
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(77) TD NO: 24:
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619-587-5360
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366.00
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                  08/617,256
                                                                         US/09/287,679
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Indels:
Gaps:
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Matches:
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Best Local Similarity:
Query Match:
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APPLICANT: K ster,
TITLE OF INVENTION:
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TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TELEFAX: 217
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TELEFAX: 217
                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
ADDRESSEE: The Square, 7th Floor
                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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APPLICATION NUMBER: 08/40
APPLICATION NUMBER: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: GEOGRAPHICA
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                             COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                        COUNTRY:
                                                                                                                        CITY: La Jolla
STATE: CA
              SOFTWARE:
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STRANDEDNESS: single
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ARE: PatentIn Release #1.0, Version APPLICATION DATA:
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(617)227-5941
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366.00
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96.43%
23.03%
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DNA Diagnostics
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Conservative:
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US-09-287-681-24
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                                                                                                                                                                Sequence 24, Application US/0928768 Patent No. 6277573
                                                      GENERAL INFORMATION:
APPLICANT: K ster, Hubert
TITLE OF INVENTION: DNA Diagnostics Based on
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White
STREET: 4250 Executive Square, 7th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
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LENGTH: 252 base pairs
TYPE: nucleic acid
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEFAX: 610-607-607
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FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 18-MAR-1996 PRIOR APPLICATION DATA:
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TOPOLOGY: linear
COUNTRY: USA
ZIP: 92037-9103
                               STATE:
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REGISTRATION NUMBER: 33,77
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                          La Jolla
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Matches:
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Indels:
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Query Match:
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Best Local Similarity:
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TELEPHONE: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/6
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                NO.:
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TOPOLOGY: lir
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TELEFAX: 619-587-5360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                GAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGCAAGCTGCGTAAGCGGCTCCTCCGC
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                                                           GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGCCCGCGAGGGCCGC
                                                                        AspAlaAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla
 GAGCGCGGCCTC
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-DB-Published_Applications_NA -QFWT|fastap -SUFFIX-p2n.rnpb -MINNATCH=0.1
-DOPCL-0 -LOOPEXT=0 -UNITS-bits -START-1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST-45 -DOCALIGN=200 -TRH_CSCORE-pct -THR_MXS-100
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-MAXLEN=200000000 -USER-US09827854_@CGN_1_1_298_@runat_11032003_101612_27578
-NCPU-6 -ICPU-3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREMSD=1 -XGAPEXT=0.5 -FGAPOP=6
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Sequence 11, Appl
Sequence 129, App
Sequence 17, Appl
Sequence 8, Appli
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US-09-827-854-11
                                 Pred. No.:
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                                                                                                                                                               SOFTWARE:
                                                                                                                TYPE: DNA
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US-10-044-090-454
US-09-827-854-12
US-09-827-854-9
US-09-954-456-760
US-09-980-107-2244
US-09-980-107-2244
US-09-980-107-244
US-09-960-352-302-133
US-09-960-352-5420
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130, Appli

131, Appli

134, Appli

135, Appli

1361, App
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/579,088
PRIOR APPLICATION NUMBER: US 09/579,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
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LENGTH: 1156
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                                          SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                        ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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Application

US/09870759

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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR FILE REFERENCE: 870759
CURRENT APPLICATION NUMBER: US/09/870,759
CURRENT FILING DATE: 2002-01-14
PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 166
SOFTWARE: PATENTIAN PATENTIAN OF SEQ ID NO 129
LENGTH: 1156
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapie
FEATURE:
NAME/KEY: CDS
LOCATION: (61)..(101)
OTHER INFORMATION:
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GENERAL INFORMATION:
APPLICANT: TERMAN, David
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RESULT 3
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Sequence 17, Applicat:
Publication No. US200:
GENERAL INFORMATION:
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SEQ ID NO 17
LENGTH: 1156
                                                                                                                                                                                                                                                                                                          LENGTH: 1156
TYPE: DNA
ORGANISM: Homo sapie
FEATURE:
NAME/KEY: CDS
LOCATION: (61)...(10
OTHER INFORMATION: NO
OTHER INFORMATION: N
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APPLICANT: Bonsal Aruna
APPLICANT: Kleyn Patrick
TITLE OF INVENTION: GENES AND POLYMORPHISMS
TITLE OF INVENTION: CARDIOVASCULAR DISEASE
FILE REFERENCE: 24735-2048
                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/802,640
CURRENT FILING DATE: 2001-03-09
NUMBER OF SEQ ID NOS: 122
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APPLICANT: Kypreos, Kyriakos E.
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TYPE: DNA
ORGANISM: Homo
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Patent No. US20020137081A1
GENERAL INFORMATION:
APPLICANY: 01ga Bandman
TITLE OF INVENTION: GENES DIFFERENTIALLY EXP
FILE REFERENCE: PA-0028 US
CURRENT APPLICATION UNMBER: US/10/044,090
CURRENT FILING DATE: 2002-01-09
NUMBER OF SEQ ID NOS: 850
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US-10-044-090-454
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TYPE: DNA
ORGANISM:
FEATURE:
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                            ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods fo
TITLE OF INVENTION: Cohlesterol levels wit
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 1156
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Best Local Similarity:
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GENERAL INFORMATION:
APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriako
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                    SOFTWARE: Fast
SEQ ID NO 9
LENGTH: 1156
TYPE: DNA
ORGANISM: Homo
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GENERAL INFORMATION:
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                                                                                                                              APPLICANT: Zannis, Vassilis
APPLICANT: Kyprecs, Kyriakos E.
APPLICANT: Kyprecs, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
              PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
                                                                                   CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
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Percent Similarity:
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                                                 LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
                                                                                                                    SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
                                                                                   ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
                                                                                                                                                                                       CAGGCCGGGGCCCGAGGGGCGCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
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PRIOR PRIOR FILLING DATE: 2000-09-25
PRIOR FILLING DATE: 2000-09-25
PRIOR FILLING DATE: 2000-09-25
PRIOR FILLING DATE: 2000-09-26
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,638
PRIOR APPLICATION NUMBER: US/60/235,711
PRIOR FILLING DATE: 2000-09-26
PRIOR FILLING DATE: 2000-09-27
PRIOR FILLING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US/60/235,720
PRIOR APPLICATION NUMBER: US/60/235,720
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PRIOR APPLICATION NUMBER: US/60/235,863
PRIOR FILLING DATE: 2000-09-27
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; ORGANISM: HOMO
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Patent No. US200:
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APPLICANT: HOTHE, DATCI T.
APPLICANT: VOCKLEY, JOSEPH G.
APPLICANT: VOCKLEY, JOSEPH G.
APPLICANT: SCHETÉ, UWE
APPLICANT: SCHETÉ, UWE
APPLICANT: SCHETÉ, INC.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 2244
LENGTH: 1157
TYPE: DNA
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                                                 ORGANISM: Homo sapi
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                                                 ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
                                                                    SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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                              GTGCAGGCTGCCGTGGGCACCAGCGCCCCCTGTGCCCAGCGACAATCAC
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score:
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CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alignment Scores: Pred. No.:
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without induc.
FILE REFERENCE: 07180/004003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 7
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{\tt ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnProl}
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CURRENT APPLICATION NUMBER: US/09/827,854

CURRENT FILING DATE: 2001-04-05

PRIOR APPLICATION NUMBER: US 09/679,088

PRIOR FILING DATE: 2000-10-04

PRIOR APPLICATION NUMBER: US 09/544,386

PRIOR FILING DATE: 2000-04-06

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                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 10
LENGTH: 1156
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without inducing hypertriglyceridemia
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ORGANISM: Homo
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138 yGluValGlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHi 158 	118 aGlnAlaArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGl 138 	98 uGlnLeuThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAl 118 	79 -ArgalaLeuMetaspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGl 98 	21036 CCCCATCCCAGCCCTTCTCCCCGCCTCCCACTGTGCGACACCCCTCCCGGCCCTCTCGGCCG 21095	78 78	6 GCCTCTGCCCCGTTCCTTCTCCCCTCTTGGGTCTCTGGCTCATCCCCCATCTCGCCCG 2	2091b TGTCTCTGCCTCTGCCTCTGCATCTGCTCTGCATCTGTCTCTGTCTCTCTC	78 78	20856 CCTCCCAAAGTGCTGGGATTAGAGGCATGAGCCACCTTGCCCGGCCTCCTAGCTCCTTCT 20915	78 78	20796 CTCACTGTGTTGCCCAGGCTGGTCTTGAACTTCTGGGCTCAAGCGATCCTCCCGCCTCGG 20855	78 78	20736 CGTCCTGGCTCTGTCTCTGTCCTTCCCTAGCTCTTTTATATAGAGAGAG	78 78	20676 TGGAATTCTCTCTCAGCTTTGTCTCTCTCTCTTCCTTTCGACTCAGTCTCTCACACT 20735	78 78	20616 CCTGGGTCTCTGCTGGTTCTAGCTTCCTCTTCCCATTTCTGACTCCTGGCTTTAGCTCTC 20675	78 78	20556 GGCGGCTATACCTCCCCAGGTCCAGGTTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGGG 20615	78 78	72 SerGlnValThrGlnGluLeu 78	b INGGATTACCTGCGCTGGGTGCAGACACTGTCTGAGCAGGTGCAGGAGGAGGAGCTGCTCAGC 2	52 TrpAspTyrLeuArgTrpValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSer 71	32 LeuArgGlnGlnThrGluTrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPhe 51	12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProGlu 31 	·09-827-854-18 (1-317) x US-09-967-013-5 (1-41907)	st Local Similarity: 60.00% Mismatches: 6 2ry Match: 86.12% Indels: 194 : Gaps: 1

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CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver. 2.0
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US-09-925-302-133
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                                                                                                                                                                                                                                                                              SEQ ID
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ORGANISM: Homo
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                                                 CGCCTGGTGCAGTACCGCGGCGAGGTGCASGCCATGCTCGGCCAGAGCACCGAGGAGCTG
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Query Match:
                                             US-09-827-854-18 (1-317) x US-09-964-824A-374 (1-478)
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CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
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APPLICANT: Horrigan, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 583
SOFTWARE: PatentIn version
SEQ ID NO 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 374, Application US/09964824A Patent No. US20020102531A1
                                                                                                                                                                                                              NAME/KEY: misc_feature LOCATION: (1)...(478) OTHER INFORMATION: n=a
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                                                                                                                                                                                                                                                                                           TYPE: DNA
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ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: Genbank Activation
NAME/KEY: unsure
LOCATION: (1)..(478)
OTHER INFORMATION: n = a or (US-09-880-107-2491
                                                                                                                                                                                                                                                                                            APPLICANT: VOCKLEY, JOSEPH G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2491
LENGTH: 478
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Patent No. US20020142981A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -09-880-107-2491/c
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-DB=EST -QFMT=fastap -SUFFIX-p2n.rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -ENDS-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DCCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-UTFEMT=pto -NORM-ext -HEARSIZE=500 -MINLEN=0 -MAXLEN=200000000
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-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAFOP=10 -XGAPEXT=0.5 -FGAPEXT=7
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Perfect score:
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3980.771 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

C 4410 442 443	C 329 303333333333333333333333333333333333	27.77.77.77.77.77.77.77.77.77.77.77.77.7	Result
107 075. 107 107 074. 070. 107 069.	1097.55 1097.55 1097 1096 1091 1091 1091 1089 1084.5 1083.5	124 124 124 121 121 121 121 121 121 121	Score
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1100 914 748 930 812 688 803	797 796 796 797 797 799 798 891	1027 1027 1042 933 933 934 942 800 922 817 817 817 817 790 770 707 741 1110 808 808 808 808 808 808 808 808 8	Length
113 113 113 113 113	13 13 13 13 13 13	13 13 13 13 13 13 13 13 13 13 13 13 13 1	DB
M91438 I60352 I55308 F96785 G76996 G70936	G71536 G71536 I15975 M04215 G70775 I60127 I55106 I55106 I55181 I54929 I45835		ΙD
114382 A 114382 A 103523 6 53085 6 67857 6 69968 6 44886 6	G715366 G7159757 G7159757 G7072753 G707750	670359 6 8 603558 6 8 8 8 9 6 7 8 8 8 8 9 8 8 8 8 8 8 8 8 8 8 8 8 8 8	Description

## ALIGNMENTS

TITLE	REFERENCE	SOURCE ORGANISM	KEYWORDS	VERSION	ACCESSION	LOCUS DEFINITION	RESULT 1 BI670367
NTH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)	<pre>Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1027)</pre>	human Homo sapiens	EST.	BI670367.1 GI:15584600	mRNA sequence. BI670367	B1670367 1027 bp mRNA linear EST 12-SEP-2001 603292738F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5312024 5',	

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DB:
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GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                         GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                                                                                                                                                                                                                GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                                                                                                                                                                                                                                                                                                                   MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
                                                    ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                                                                               LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                         CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11790 row: 1 column: 09
High quality sequence stop: 845.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tissue Procurement: Miklos Palkovits, M.D., Ph.D. cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki Toshiyuki and Piero Carninci (RIKEN) cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1999)
Contact: Robert St
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Email: cgapbs-r@mail.nih.gov
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/db_xref="taxon:9606"
/clone="IMAGE:5312024"
/clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
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AUTHORS
TITLE
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BG763371.1
EST.
                                                                                                                        cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov bcalate: LLCM1718 row: b column: 10 High quality sequence.stop: 817.
                                                                                                                                                                                                                                                    Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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/lab_host="DH10B (phage-resistant)
                                                               /organism="Homo sapiens"
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                                                                                                           ocation/Qualifiers
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Best Local Similarity:
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                                                               LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMet-Gl
                                                                                                              ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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                                                  CTACAGGAGCGGGCCCAGGCCTGGGGGCGAGCGGCTGCGCGCGGGATGGAGGAGATGCGG
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GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40
                                                GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGAGACCGAGTGGCAGAGC
                                                                                                           ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
Toshiyuki and Piero Carninci (RIKEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, Mammalian
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                      1 (bases 1 to 938)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mamme
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
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cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (1 DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information
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http://image.llnl.gov
Plate: LLCM1674 row: c colu
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/db_xref="taxon:9606"
/clone="MAGE:484141"
/clone="NHH_MGC_49"
/tissue_type="melanotic melanoma, high MDR (cell line)"
/tosue_type="melanotic melanoma, high MDR (cell line)"
/tosue_type="melanoma, high MDR (cell line)"
/tosue_type="melano
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Email: cgapbs r@mail.nih.gov

Tissue Procurement: Miklos Palkovits, M.D., Ph.D.

cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carhinci (RIKEN)

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
platto./limage.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 942)
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603249241F1 NIH_MGC_96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plate: LLAM11762 row: a column:
High quality sequence stop: 762.
Location/Qualifiers
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                                                                                                                                    177
                                                                                                                                                                                                                                  /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: Sal1-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for the primary enriched for the primary library enriche
                                                                                                                               for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

284 c 36i g 120 t
                                                                                                                                                                                                                                                                                                                                               /tissue_type="hypothalamus"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5301010"
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                                                                                                                                                                                                                                                                                     GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                    ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
                                                                                                            LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                                                                                                                                                                                                                                                                                                                                      GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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  ACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
                                                                                     CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
                                                                                                                                                                           CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAGGTCACCCAGGAACTGAGGGCG
                                                                                                                                                                                                                                                                                                                                                 GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCAGAGCCGAGTGGCAGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov plate: LLCM1875 row: i column: 03 High quality sequence stop: 792. Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
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Clone distribution: MGC clone distribution information can
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
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/clone="IMAGE:5420618"
/clone=lib="NH_MGC_112"
/tissue_type="melanotic melanoma, cell line"
/tissue_type="melanoma, cell line"
/tissue
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Primates;
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Mismatches:
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Gaps:
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ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro|
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.
Plate: LLCM1419 row: k column: 08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA sequence.
BG472299
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1 (bases 1 to 927)
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              170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    quality sequence stop: 848
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/tissue_type="retinoblastoma"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
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/note="Organ: eye; Vector: potB7; Site_1: xhoI; Site_2:
/lab_host="Note: the following 5"
cloned into EccRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Library constructed by Ling Hong
in the laboratory of Gerald M. Rubin (University of
Callifornia, BerKeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
70 a 271 c 371 g 115 t
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                       CCGGCCTCAAGAGCTGGTTCGGAGCCCTGGGTGGAAGACATG
                                                  aArg-LeuLysSerTrpPhe-GluProLeuValGluAspMet 290
                                                                                         gAlaLysLeuGluGluGlnAlaGlnGİnIleArgLeuGlnAlaGluAla---PheGlnAl
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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                                                                                       GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                   CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI)
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov Plate: LLAM11760 row: b column: 04
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National Institutes of Health, Ma
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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Mammalia; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

273 c 359 g 114 t
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/lab_host="DH10B"
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/clone_lib="NIH_MGC_96"
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/db_xref="taxon:9606"
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BG774871.1 GI
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Eukaryota; Me
Mammalia; Eut
                                                                                                                                                                Contact: Robert Strausberg, Ph
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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602649975F1 NIH_MGC_40
                                                                                found through the I.M.A.G.E. Con-
http://Image.llni.gov
Plate: LLCM1612 row: e column:
High quality sequence stop: 813.
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                                                                                                                                                                                                                                                                     human.
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4760827"
/clone_lib="NIH_MGC_40"
/tissue_type="carcinoma, cell line
/lab_host="DH10B (phage-resistant)
                                                                       Location/Qualifiers
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Primates;
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                                               LeuGlnGluArgAlaGlnAla--TrpGlyGluArgLeuArgAlaArgMetGluGluMetG
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                                                                                    CCCCTGGTGGAACAGGGCCGCGTGCGGGCCTGCACTGTGGGCTCCCTGGCCGGCAG-CCG
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Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of Callifornia, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).

Note: this is a NIH_MCC_Library."
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MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11726 row: g column: 10
High quality sequence stop: 783.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausperg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 811)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: Sal1-xhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched
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/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:5287329"
/clone_lib="NIH_MGC_96"
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                                                                                                   cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information car found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1790 row: g column: 17 High quality sequence stop: 829.
                                                                                                                                                                                                               Unpublished (1999)
Contact: Robert Strausberg, E
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, P
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/organism="Homo sapiens"
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                                                                                                                      GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                                                                                                                                                                                                                                                 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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CTACAGGAGCGGGCCAGGCTG---GGGCGAGCGGTGCGCGCGCGCGATGGAGGAGGATGGGC
                                                                                                                                                                                CAGGCCGGGGCCCGAGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
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Directionally cloned into ECORI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "

a 244 c 342 g 106 t
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밁 δÃ 밁 ð Ъ δÃ В Qy В δõ 뫄 δ 뫄 δõ Дb Qy В δÃ B δ В δÃ В QΥ Дb Qy

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BASE COUNT
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AUTHORS
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.
CDNA Library Preparation: Michael J. Brownstel
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consor
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution infi
found through the I.M.A.G.E. Consortium/LLNL at
http://image.llnl.gov
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BG707147.1 GI
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Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, M
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Mammalia; Eutheria;
1 (bases 1 to 790)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4793084"
/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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                                                                                                                                             NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Ge
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D.,
CDNA Library Preparation: Michael J. Brown
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Co
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                                                                               DNA Sequencing by: Incyté Genomics, Inc.
Clone distribution: MGC clone distribution i
found through the I.M.A.G.E. Consortium/LLNL
http://image.llnl.gov
plate: LLAM11691 row: 1
High quality sequence stop: 812.
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603194314F1 NIH_MGC_95
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 919)
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Location/Qualifiers
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Brownstein
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/note="Organ: brain; Vector: pBluescriptR (modified pBluescript RS+); Site\_1: BanHI; Site\_2: SalI-xhoI (gtcgag pBluescript RS+); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.5 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH\_MGC Library."

Percent Similarity: Best Local Similarity: US-09-827-854-18 (1-317) x BI551475 (1-919) GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly LysLeuCysLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg CAGCCGGACCCGGGACG - - CTGGACGAAGTGAAGAGCACGTGGGCGGAGGG - TGCGCCAA ySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTTGGGATTACCTGCGCTGGGTGCAGACA GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGACCGAGTGGCAGAGC CAGGCCGGGGGCCGCGATGGCGCGCGAGCGCGTGCTCAGCGCCATCCGCGAGCGCCTGGGG AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC 1.37e-109 1141.00 93.26% 92.13% 71.81% Indels: Gaps: Length: Matches: Conservative: Mismatches: 919 246 3 17 4 0 140 847 260 790 240 220 672 200 612 180 552 160 492 432 120 372 100 312 80 252 60 40

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               GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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Tissue Procurement: DCTD/DTP
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National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="melanotic melanoma, cell line"
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/note="Organ: skin; Vector: pOTBF; Site_1: XhoI; Site_2:
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/note="Organ: skin; General delay adaptor: good of the company of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Libbrary."

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/clone_lib="NIH_MGC_112"
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                                                                                                                                                                                                                                                       Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgpabs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
                                                                                                                                                                                               Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov plate: LLAM10669 row: i column: 23
                                                                                                                                      High quality sequence stop: 832.
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National Institutes of Health, Mammalian
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            /tissue_type="hypothalamus"
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                                         /db_xref="taxon:9606"
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                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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Brownstein
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US-09-827-854-18 (1-317) x BG706129 Query Match: DB: Best Local Alignment Scores: BASE COUNT Percent Similarity: No.: 792 675 615 495 435 375 315 180 555 160 120 101 255 195 81 61 41 21 75 gLysLeuCysLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTy aArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVa ThrProValAlaGluGluThrArgAlaArgLeuSerLysGlu-LeuGlnAlaAlaGlnAl MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal CCAGGCCGGGGCCCGAGGGGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGG rGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeu-G lGlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuAr CCGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGT GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40 GCAG-CGGACCCGG----ACGCTGGACGAGTG-AAGGAGCAGTGGGGAAGTGTGCGC lySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArg 258 CAAGCTGCGTAAGCGGCTCCTCCGCGATGCCCGATGACCTGCAGAAGCGCCTGGCAGTGTA GCAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCG GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA GAGCAAGCGGTGGAGACAGAGCCGGAGCCGGAGCTGCGCAGCAGACCGAGTGGCAGAGC ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG Similarity: , 171 ۵ pBluescript Ks+); Site\_1: BamHI; Site\_2: SalI-XhOI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH\_MGC Library."

a 260 c 358 g 117 t 1 others 1.5e-107 1121.50 94.23% 93.46% 70.58% (1-907)Length:
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-Q=/Cgn2_1/USPT0_spool/US09827854/runat_11032003_101610_27486/app_query.fasta_1.3576
-DB=GenEmbl -QFMT=fastap -SUFFIX=p2n.rge -MINMATCH=0.1 -LOOPCL=0 -LOOPEXY=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
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-USER=US09827854 eCGN_1_1_13008_@runat_11032003_101610_27486 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
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36: em_htg_vrt:*
37: em_htg_vrt:*
38: em_sy:*
40: em_htgo_hum:*
41: em_htgo_other:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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AR164342 AR205885 S76779 MUSAPOEE RATAPOE AC127479 AC073760 MUSAPE AR164387	E00823 AX302545 BD004278 BD004278 BD004277 AX333278 BC003557 BD004279 BD004279 BD004279 HUMAPOE BD004279 HIMAPOE AF261279 AX358722 AF050154 AC011481 HUMAPOE4 AF261280 BABAPOE4 AF261280 BABAPOE4 AF261280 BABAPOE4 AF303833 BTAPOLPE AX384545 AX384545 BYASEE AX384541 AX384591 AX	ID
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PF 29-NOV-198
PF TERANISHI
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strandedness: Double;
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                                                                                                                                                                          ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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/organism="unidentified"
/db_xref="taxon:32644"
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/product='human
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/db_xref="taxon:9606"
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                                                                                   Patent: JP 2001017028-A 2 2
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/2
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128
PR SHINOBU FUJITA, HIROKI
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                                                          PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE A01K67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC C12N15/09//C07K14/775, PC (C12N5/10.C12P1.C1.
                                                                                                                                                                Eukaryota; Metazoa; Chordata; Cre
Mammalla; Eutheria; Primates; Cat
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Fujita,S., Hamanaka,H., Fukui,Y.
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          Location/Qualifiers (61)..(1011).
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/organism="Homo sapiens"
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                                          CTGGAGGAGCAGGCCCAGCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCCCGCCTCAAG
                                                                             AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCGCCAAG
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J. Clin.
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84212473
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85279526
                                                                                                                                                                                                                                                                                                                                   5 (bases 577 to 624)
Gill,L.L., Peoples,O.P., Pearston,D.H., Robertson,F.W.,
Humphries,S.E., Cumming,A.M. and Hardman,N.
Isolation and characterisation of a variant allele of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 (bases 250 to 777) Wallis, S.C., Rogne, S., Gill, L., Markham, A., Edge, M., Woods, D. Williamson, R. and Humphries, S. The isolation of cDNA clones for human apolipoprotein E and detection of apoE RNA in hepatic and extra-hepatic tissues EMBO, J. 2 (12), 2369-2373 (1983)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  J. Biol.
83082756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 355 to 1156) Breslow, J.L., McPherson, J., Nussbaum, A.L., Williams, H.W., Lofquist-Kahl, F., Karathanasis, S.K. and Zannis, V.I.
                                                                                                                                                                                             Type III hyperlipoproteinemia associated with apolipoprotein phenotype E3/3. Structure and genetics of an apolipoprotein is
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human apolipoprotein E of a new variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    McLean, J.W.,
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Synthesis, intracellular
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Homo sapiens preapolipoprotein E
[6] sites; mutations resulting in type III hyperlipoproteinemia. Apo E is a component of normal human very low density lipoprotein. There are six human apo E phenotypes known to result from a single structural gene, three of the common alleles being epsilon-4,
                                                                 epsilon-3 and variant. [5] epsilon-2 allele.
                                                                                                                                                                                                                           Rall,S.C. Jr., Newhouse,Y.M., Clarke,H.R., McCarthy,B.J., Mahley,R.W. and Bersot,T.P.
                                                                                                                                                                                                                                                                                                          human apolipoprotein E
Biochem. Biophys. Res.
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 88
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Elshourbagy, N.A., Chang, D.J., Mahley, R.W. and
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J. Biol. Chem. 258, 11422-11422 (1983)] is too extensively revised relative to [1] to record the revisions in the FEATURES table. The sequence below is that of [J. Biol. Chem. 258, 11422-11422 (1983)]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is thought to result from a c to t change (arg to cys) at base 586 below [3],[5]. The sequence shown is 578 homologous with human apo A-I and 818 homologous with rat apo E. For the epsilon-4 sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                epsilon-3 and epsilon-2. This sequence appears to be of the epsilon-3 allele. [1] argues that the apo E polymorphism involves mutations in the structural coding region; for example the epsilon-2 phenotype which is characterized by hyperlipoproteinemia
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/citation=[4]
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ELALGRFWDYLRWVQTLSEQVQEELLSSQVTQELRALMDETMKELKAYKSELEEQLTP
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/db_xref="GI:178851"
/db_xref="GDB:G00-119-691"
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                                                                               GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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586
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448
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Direct Submission
Submitted (20-FEB-2001) National Institutes of Submitted (20-FEB-2001), Cancer Genomics Office, Gene Collection (MGC), Cancer Genomics Office, Room 11A03, Betheso
                                                                                                                                                                                                                                                                        Homo sapiens,
complete cds.
BC003557
BC003557.1 GI
                    cDNA Library Preparation: Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. CODNA Sequencing by: Genome Sequence Centre, BC Cancer Agency, Vancouver, BC, Canada
                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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Clone distribution: MGC clone distribution information can be fo
through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
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VAEETRARLSKELQAQARLGADMEDVCGRLVQYGEVQAMLGGTEELRVRLASHLR
KLRKRLLRDADDLQKRLAVYQQGARREGAERGLSAIRERGLPLVEOGRVVRAATVCSLAG
QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
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/protein_id="AAH03557.1"
/db_xref="G1:]1907699"
/translation="MKVLWAALLVTFLAGCQAKVEQAVETEPEPELRQQTEWQSGQRW
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/clone="MGC:1571 IMAGE:3355712"
/tissue_type="Eye, retinoblastoma"
/clone_lib="NIH_MGC_16"
/lab_host="DHi0B-R"
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/db_xref="LocusID:348"
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                      JP 1994315392-A/1.
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                                                                                                                     Homo sapiens (human)
pr 1994315392-A/1
15-NOV-1994
11-JUN-1985 JP 1994015433
MORIMOTO HIRONORI, TERANISHI YUTAKA
MORIMOTO, COTK13/00,C12N5/10,C12N15/12,C12N15/18,(C12P21/02,C12R1:91),
                              sig_peptide
mat_peptide
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                                                                               Key
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                                                                                                        (C12N5/10,C12R1:91);
/product='human
Location/Qualifiers
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                              /organism='Homo
15. .68
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                    apolipoprotein E3'
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Fujita; S., Hamanaka, H., Fukui, Y Apo E humanized mammal Patent: JP 2001017028 A 1 23-JA MITSUBISHI CHEMICAL CORP
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A01K67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC
C12N15/09/C07K14/775,
PC (C12N5/10, C12R1:91), C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
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JP 2001017028-A/1
23-JAN-2001
28-APR-2000 JP 200012
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Matches:
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Sequence 378
AX333278
AX3333278.1
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Patent:
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Mammalia; Eutheria;
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Location/Qualifiers
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Primates;
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Matches:
Conservative:
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Indels:
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Gene expression profiles in liver cancer
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/db_xref="taxon:9
/map="19q13.2"
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/gene="APOE"
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                /tissue_type="liver"
/dev_stage="57-year (
                                       /cione="pHAE[112,178,813]."
/sex="male"
/tio--
                                                                                         Location/Qualifiers
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                                                                             CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                     LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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/note="t in preplace="c"
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/note="a in p
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VAEETRARLSKELQTAQARLGADMEDVCGRLVQYRGEVQAMLGQSTEELRVRLASHLR
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/replace="a"
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QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
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/note="G00-119-691"
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REFERENCE
AUTHORS
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PN JP 2001017028-A/3
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128
PR SHINOBU FUJITA, HIROKI
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        PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE YOKOYAMJAOLIKOT/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC C12N15/09/C07K14/775, PC (C12N5/10, C12R1:91), C12N5/00, C12N15/00, (C12N5/00, C12R1:91)
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Mammalia; Eutheria;
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                           ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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Location/Qualifiers
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/db_xref="taxon:9606"
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300 100 100 360 120 120 40 180 20

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		GOPW OLSP SHLR SLAS EAFO		13,	,. ,.	1995		
RESULT 15 AF261279 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM		Qy 2 Db 8 Qy 2 Db 8	Oy 2 6	Oy 16 Db 56 Oy 18 Db 62	Qy 1 Db 4 Qy 1 Db 5	Qy Db 3 Qy 1 Db 3	Qy Db 2 Qy Db 2	Db Qу Db 1
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PUBMED
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AUTHORS
TITLE
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1. .5491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nickerson, D.A., Taylor, S.L., Fullerton, S.M., Weiss, K.M., Clark, A.G., Stengard, J.H., Salomaa, V., Boerwinkle, E. and Sing, C.F. Sequence diversity and large-scale typing of SNPs in the human
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Nickerson, D.A.
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Genome Res. 10 (10), 1532-1545 (2000)
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163
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2440
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118 3902	uThrargAlaargLeuSerLysGluLeuGlnalaal 	98 uGlnLeuThrProValAlaGluGluTh 	Оу ( Db 38
98 3842	tLysGluLeuLysAlaTyrLysSerGluLeuGluGl 	79 -ArgAlaLeumetAspGluThrmet 	Qy
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3242	TGAGTGTCCCCATCCTGGCCCTTGACCCTCCTGGTG	183 TCCCAGGTCACCCAGGAACTGAGG	Db 31
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71 3182	31nThrLeuSerGluGlnValGlnGluGluLeuLeuSer 	52 TrpAspTyrLeuArgTrpValo	Qу Db 31:
3122	GCCAGCGCTGGGAACTGGCACTGGGTCGC	63 CTGCGCCAGCAGACCGAGTGGC	Db 30
51	lnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPhe	32 LeuArgGlnGlnThrGluTrpGlr	Qy
31 3062	ValGluGlnAlaValGluThrGluProGluProGlu 	12 PheLeuAlaGlyCysGlnAlaLys	Qy :
	9 (1-5491)	827-854-19 (1-317) x AF26127	US-09-8
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Оу :	Qy :	Οy :	Qy :	Qy Db 4:	Qy Db 4	Qy Db 4	Qy Db 4	Qy 3	Db 3
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4443	4383	4323	4263	4203	4143	4083	4023	3963	
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Search completed: March 14, 2003, 17:32:49
Job time: 2313.23 secs

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## SUMMARIES

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_	Human EST-derived	Human secreted pro	Human apolipoprote	Human diagnostic a	Thyroid cancer rel	Colon adenocarcino	Gene #2492 used to	Human cDNA for nov	()	Lung cancer associ	Human cDNA encodin	ApoE4L protease cD	ס	Partial human apol	Human ApoE4Lx2 cDN	ApoE4Lx2 protease			alpha	O)	apolip	Apo E	-	apoli	_					Human ApoE4 coding		cancer :	n heni	Gene #2244 ised to			-	aport	-	polidod	numan aportpoprote			- 1	Description

## ALIGNMENTS

RESULT 1 AAD22052 CDS Human apolipoprotein E (apoE) allele, apoE2\*\* DNA Human; apolipoprotein E; apoE; cholesterol; atherosclerosis; hypertriglyceridaemia; low density lipoprotein; LDL; ds. 12-FEB-2002 AAD22052 standard; DNA; 1156 BP Homo AAD22052; sapiens (first entry) /\*tag= Location/Qualifiers

FFT XXX

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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04-OCT-2000;
05-APR-2001;
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                                                                                                                                                                                                                                                                                   drugs for treating diseases associated with APDE activity, e.g. familial dysbetalipoproteinaemia, type III hyperlipoproteinaemia, atherosclerosis, and Alzheimer's disease. They are useful to validate APDE as a candidate agent for treating a specific condition or disease predicted to be associated with APDE activity and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with APDE activity. Genotyping or haplotyping methods are useful to screen for compounds targeting APDE to treat a specific condition or disease associated with APDE activity. The present sequence is a CDNA encoding human APDE protein.

APDE gene is located on chromosome 19q13.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                E (APOE) gene. The invention also relates to compositions and met for haplotyping and/or genotyping the APOE gene. The haplotyping methods of the invention are useful for improving the efficacy are reliability of several steps in the discovery and development of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 26; Fig 2; 78pp; English
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                        GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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DB; AAE15158.
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GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                             LysLeuArgGlnArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
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                                                            CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
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                                                                                                                   Detecting and identifying ovarian tumor, identifying increased risk for developing ovarian cancer, and determining effectiveness of ovarian cancer treatment, by measuring expression level of ovarian tumor marker
                                                 Claim
                                                                                               gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mucinous cystadenoma; borderline mucinous tumour; endometrioid carcinoma undifferentiated carcinoma; clear cell adenocarcinoma; cystadenofibroma; adenofibroma; Brenner tumour; serial analysis of gene expression, SAGE; immune response pathway; cell proliferation regulation; protein folding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ovarian tumour marker gene; human; overexpression; upregulatiepithelial tumour; cancer; diagnosis; prognosis; disease monidentification; serous cystadenoma; borderline serous tumour; serous cystadenocarcinoma; mucinous cystadenocarcinoma;
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membrane localised; secreted; therapeutic target; cytostatic;
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Best Local Similari
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prognostic markers, the ovarian tumour marker genes or their encoded proteins may be used as therapeutic targets for the treatment and prevention of ovarian cancer. Sequences ABA83081-ABA83122. ABA83180, ABA83182 and ABA83184 represent the ovarian tumour marker genes of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tumours in an individual via the detection and measurement of the expression of ovarian tumour marker genes (ABA83081-ABA83122, ABA83180, ABA83181), and ABA83184) or segments thereof (ABA83123-ABA83169, ABA83179, ABA83181 and ABA83183). The methods of the invention are useful for detecting an ovarian tumour in a patient, for identifying an individual at increased risk for developing ovarian cancer, in prognostic tests for assessing the relative severity of ovarian cancer, in tests for
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LysLeuArgGlnArgLeuLeuArgAspA|aAspAspLeuGlnLysArgLeuAlaValTyr
                                LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGlnGlnLeu
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The present invention relates to an ApoE humanised mammalian cell. present sequence is the coding sequence for human ApoE3, which was in the method of the present invention. The ApoE humanised mammalia can be used for screening for agents useful for treating or prevent

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Matches:
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The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (Li receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-APR-2000;
04-OCT-2000;
05-APR-2001;
                                                                                                                                                                                                                                                                                                                       New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia -
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(UYBO-)
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hypertriglyceridaemia; low density lipoprotein;
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                                                                                                         LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
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                                                                                                                                                                                                                                   Percent Similarity:
                                                                                                                                                                                                                                                                                          present sequence is the coding sequence for in the method of the present invention. The can be used for screening for agents useful Alzheimer's disease and arteriosclerosis.
                                                                                                                                                                                                                                                                                                                                                         useful for treating arteriosclerosis -
                                                                                                                                                                                                                                                                           Sequence 1156
                                                                                                                                                                                                                                                                                                                          The present invention relates to
                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 11-12;
                                                                                                                                                                                                                                                                                                                                                                New apoE humanized useful for treating
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ApoE humanised mammalian ce
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1156
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New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in mammal, without inducing hypertriglyceridemia -
English.
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The present sequence is a human apolipoprotein E (apoE) allele, apoE2\* DNA. The apoE lipoproteins are useful for lowering toholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (Loreceptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or haddenitation of lipoprotein remnants in the bloodstream or haddenitatin the bloodstream or haddenitation or haddenitation or haddenitat defect in remnant removal. or having (LDL)

Sequence 1156 BP; 208 A; 367 Ç; 432 <u>ი</u> 149 T; 0 other;

3.48e-110 1573.00 99.68% 99.37% 99.24%

Length:
Matches:
Conservative:
Mismatches:
Indels:

1156 315 1 0 0

Gaps:

US-09-827-854-19 (1-317) x AAD22051 (1-1-156) 120 20

GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC 60 180 40

GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA 80 240

100 300

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ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal

GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG

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LysLeuArgGlnArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCCACCTGCGC 180 540

GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly CAGGCCGGGGCCCGAGGGCGCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG AAGCTGTGTAAGCGGCTCCTCCGCGATGCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC 660 200 600

ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 240 220

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by preparing a genetic construct (contg. an ApoE-encoding sequence, e.g. AAT06957, and flanking regulatory sequences enabling the protein to be expressed in insect cells), which is then introduced into a Manduca sexta larva (using a recombinant Autographica californica nuclear polyhedrosis virus) and recovering the protein from the haemolymph of the larval host. The ApoE produced is in a form sufficiently complexed with lipids to be biologically active, which cannot be achieved in insect cell cultures, and can therefore be used in therefore be
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nuclear
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haemolymph; li
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                                     LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys
                                                           SerargThrArgAspArgLeuAspGluValLySGluGlnValAlaGluValArgAlaLyS
                                                                                                  LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
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                                                                                                                                                                                                                                                                                                                                                                                 expression of two or more genes represented in ABN93503-ABN97455 in a tissue sample. The method of the invention has hepatotropic, and cytostatic activity. The method is useful for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma and metastatic liver carcinoma in a patient. The method is useful for identifying expression profiles which serve as useful diagnostic markers as well as markers that can be used to monitor disease states, disease progression, drug toxicity, drug efficacy and drug metabolism.

Note: The sequence data for this patent did not form part of the printed as pecification, but was obtained in electronic format directly from WIPO at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the printed at the 
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                                                                                                                                                                                                                                                                                                                                                                      at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel method for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma or metastatic liver tumour in a patient, and differentiating metastatic liver cancer from hepatocellular carcinoma in a patient, involving detecting the level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
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                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of diagnosing (1) the onset or CC progression of benign prostatic hyperplasia (BPH), or screening (II) for Cr dentifying an agent that modulates the onset or progression of BPH.

CC The method is based on changes in gene expression in BPH tissue isolated CC from patients exhibiting different clinical states of prostate (CC thyperplasia as compared to normal prostate tissue. (I) comprises CC detecting the expression levels of one or more genes in prostate cells (II) comprises preparing a first gene expression profile CC of BPH cells or BPH-like cell population, exposing the cells to the CC agent, preparing a second gene expression profile CC cells, and comparing the first and second gene expression profiles. (CI) is useful for diagnosing the onset or progression of BPH. (II) is useful for identifying an agent that modulates the onset or progression CC of BPH. The methods are useful to present information identifying CC level of genes given in the specification in the tissue or cells to the CC level of expression of gene in the database, and displaying the CC level of expression of gene in the database, and displaying the compared to the expression level in BPH. Agents using (II) are useful for compared to the expression level in BPH. Agents using (II) are useful for the agent prostatic burserias agenc sequences of the invention.
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05-JUN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying drugs for and diagnosing benign prostatic hyperplasia, by detecting expression levels of one or more genes in prostate cells from patient that are differentially regulated compared to normal prostate
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30-MAY-2001;

2001WO-US10838

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CC The present invention describes a method (M1) for screening for an anti-neoplastic agent. The method involves exposing cells to a chemical CC agent to be tested for anti-neoplastic activity, determining a change in CC expression of at least one gene (I) of a signature gene set, where (I) CC comprises a sequence (S) selected from 8447 sequences (given in ABL61664 Cto ABL70110), or is at least 95% identical to (S), where a change in CC expression is indicative of anti-neoplastic activity. (I) has cytostatic CC activity and can be used in gene therapy. M1 can be used for screening CC an anti-neoplastic agent, and can be used for producing a product which compared to the anti-neoplastic agent as a ccompared to the anti-neoplastic agent as a ccompared to the anti-neoplastic agent as a ccompared to concert such as colon, breast, stomach, lung, thyroid, CC cosophageal, ovarian, kidney, prostate or pancreatic cancer, infiltrating ductal cancer, infiltrating ductal cancer, infiltrating ductal cancer, and the concert, squamous cell cancer, infiltrating ductal cancer, infiltrating ductal cancer.
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05-JUN-2000

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03-OCT-2000

03-OC
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                                                                             Location/Qualifiers 61..1014
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28-APR-2000;

2000JP-0128919

220	GluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro	01 ProLeuVa 	Qy 2
660	CGGCCTCAGCGCCATCCGCGAGCGCCTGGGG	01 CAGGCCGGGGGCCCCGCGAGGGCCGC	Db 6
200	LeuGly	81 GlnAlaGlyAlaArgGluGlyA	0у 1
180	gaspalaaspaspleuglnlysargleualavaltyr 	61 LysLeuArgGlnArgLeuLeuA              1 AAGCTGCGTAAGCGGCTCCTCCC	Qy 1 Db 5
160 540	lnalaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 	81 C	Qy 1 Db 4
140 480	gLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 	21 Ar 21 CG	Qy 1 Db 4
120 420	ProValAlaGluGluThrArgAlaÅrgLeuSerLysGluLeuGlnAlaAlaGlnAla 	01 Thr     61 ACC	Oy 1 Db 3
100 360	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu 	81 01	Qy Db 3
300	LeuSerGluGlnValGlnGluGluLeuLeuLeuSerSerGlnValThrGlnGluLeuArgAla 	61 LeuSerGluGlnValGlnGluGlu 	Оу Db 2
60 240	pGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 	41 GlyGlnArgTrpGluLeuAlaLeu 	Qy Db 1
40	lnalaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 	21 GluGlnAlaValGluThrGluProv 	Оу Db 1
20 120	ValleutrpalaalaLeuLeuValthrPheLeuAlaGlyCysGlnAlaLysVal 	1 MetLysValLeuTrpAlaAlaLeu 	Оy
	6 (1-1156)	27-854-19 (1-317) x AAF8431	US-09-8
	Length: 1156 Matches: 315 Conservative: 1 Mismatches: 1 Indels: 0 Gaps: 0	nment Scores: 6.98e-110 e: 1569.00 ent Similarity: 99.68% Local Similarity: 99.37% y Match: 98.99% 22	Alignmer Pred. No Score: Percent Best Loc Query Ma DB:
	C; 432 G; 147 T; 0 other;	equence 1156 BP; 208 A; 369	S
The used an cell ting	relates to an ApoE humanised mammalian cell. The coding sequence for human ApoE4, which was use resent invention. The ApoE humanised mammalian on agents useful for treating or preventing arteriosclerosis.	present invention : sent sequence is the the method of the pube used for screen heimer's disease and	CC The CC in CC can
	Japanese.	sclosure; F	Di
	cell useful for screening for agents ting Alzheimer's disease and	w apoE humanized mammalian eful for treating or preven teriosclerosis -	0 5 7
		ºI; 2001-285406/30. ·PSDB; AAB80998.	Q. W
		(MITU ) MITSUBISHI CHEM CORP.	
		-MAY-1999; 99JP-0125647.	06

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Search completed: March 14, 2003, 12:43:25 Job time: 183.728 secs

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Perfect score:
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   1134
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998
998
980
507
370
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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/Dackfiles1 seq:*
6: /cgn2_6/ptodata/2/ina/Dackfiles1 seq:*
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Ygapop 10.0 , X
Fgapop 6.0 , E
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US-08-949-155-5
US-08-949-155-51
US-08-949-155-51
US-09-819-964-51
US-08-726-306A-28
US-07-849-389-6
US-08-617-256-24
US-09-287-141-24
US-09-431-13-24
US-09-504-245-24
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51, Appl
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  INFORMATION FOR
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ZIP: 53701-2113
                                                                                                                                                                                       CITY: Madison
STATE: WI
                   TELEPHONE:
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## ALIGNMENTS

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Sequence 1, Application US/07709949
Patent No. 5472858
Patent No. 5472858

GENERAL INFORMATION:
APPLICANT: Attle, Alan D
APPLICANT: Sturley, Stephen L
APPLICANT: Beckage, Nancy E
TITLE OF INVENTION: Insect Larvae
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSE: Quarles & Brady
STREET: P.O. Box 2113
COUNTRY: USA
ZIP: 53701-2113
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/07/709,949
FILING DATE: 19910604
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 9629691801
TELECOMMUNICATION INFORMATION:
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SEQUENCE CHARACTERISTICS: LENGTH: 1157 base pairs

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Best Local Similarity:
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NAME/KEY: misc_signal
LOCATION: 62.119
PUBLICATION INFORMATION:
AUTHORS: McLean, J W
JOURNAL: JOURNAL of Bio
VOLUME: 259
PAGES: 6498-6504
DATE: 1984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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ORIGINAL SOURCE:
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ANTI-SENSE: NO
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LOCATION:
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GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                LysLeuArgGlnArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                                                                        ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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                                                                                                              CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
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                       AAGCTGCGTAAGCGGCTCCTCCGCGATCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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Journal of Biological Chemistry
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                                                                                                                                                           APPLICATION NUMBER: US 60/027,338
FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM FOR COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Piedrahita, Jo
APPLICANT: Bazer, Fuller
TITLE OF INVENTION: Compo
TITLE OF INVENTION: Gener
                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                    STREET: P.O. BOX
CITY: Houston
STATE: TX
COUNTRY: US
ZIP: 77210-4433
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             STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                NAME: Hibler, David W. REGISTRATION NUMBER: 41,071
                                                                                                       TELEPHONE:
                                                                                                                                     REFERENCE/DOCKET NUMBER:
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5. 6271436
                                      nucleic acid
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                                                   1126 base pairs
                                                                                           : (512) 418-3000
(713) 789-2679
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Query Match:
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; LOCATION:
US-08-949-155-5
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           ArgLeuLysSerTrpPheGluProLeuVålGluAspMetGlnArgGlnTrpAlaGlyLeu
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CTCCTCAAAGGCTGGTTCGAGCCTCTGGTGGAAGACATACGGCGCCAGTGGGCCGGGCTG
                                                                                                    GAGATGGGCAGCCGGACCGCCCTGGATGAGATGCGTGAGCAGCTGGAGGAGGTG
                                                                                                                    GluMetGlySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluVal
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                                                 CGCACCAAAGTGGAGGAGCAGGCAGCCAGTTGCGCCTGCAGGCCGAGGGATTCCACGCC
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RESULT 3
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                                                                                                                                                                                                                                                            Alignment Scores:
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GENERAL INFORMATION:
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TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
 111
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                            21 GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu 37
                                                                              1 MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: <UNKNOWN>
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRANION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK:177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/819,964
FILING DATE: 28-Mar-2001
CLASSIFICATION: <Unknown>
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: ARNOLD,
                                                                                                                                                                                                                                                                                                            SEQUENCE DESCRIPTION:
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LOCATION:
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 1126 base pairs
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Matches:
Conservative:
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                                                                                                                                                                  Sequence 51, Application US/08949155 Patent No. 6271436
                                                                                                                                                       GENERAL INFORMATION:
STATE: TA
COUNTRY: US
77210-4433
                                                                               NUMBER OF SEQUENCES: 5
                                                                                                       TITLE OF INVENTION: Compositions and Methods TITLE OF INVENTION: Generation of Transgenic
                                                                                                                              APPLICANT: Piedrahita, Jorge APPLICANT: Bazer, Fuller W
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                                                          ADDRESSEE: ARNOLD, STREET: P.O. Box 44
                                               CITY: Houston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/
FILING DATE: 09-MAY-1997
ATTORNEX/AGENT INFORMATION:
                          2837
                                                                             2777
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REFERENCE/DOCKET NUMBER: TA
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US/08/949,155
FILING DATE: CONCULTENTLY Herewitl
CLASSIFICATION: 800
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MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                        2598
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                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
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                          TTGGGGGGAAAAAACTTTTTTTTTTTTTTTCATTTGACTTCATGTCTTGCTTTCCTTTCC 2896
                                                                              CCCTCCTGGCGAACCGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
                                                                                                                                                                                    TGGCAGGGCAGCCAGCCCTGGGAGCAGGCCCTTGGGGCCGCTTCTGGGATTACCTGCGCTGG
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(713) 789-2679
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998.00
58.43%
48.76%
62.97%
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Conservative:
Mismatches:
Indels:
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RESULT 5 US-09-819-964-51 ; Sequence 51, Application US/09819964 ; Patent No. 6369294 ; GENERAL INFORMATION: Piedrahita, Jorge Bazer, Fuller WITITLE OF INVENTION: Compositions and Methods for the Generation of Transgenic Animal Species ; NUMBER OF SEQUENCES: 51 ; CORRESPONDENCE ADDRESS: ; ADDRESSE: ARNOLD, WHITE AND DURKEE ; ADDRESSE: ARNOLD, WHITE AND DURKEE ; TREET: P.O. Box 4433 ; CITY: Houston	Qy 293 GlnTrpAlaGlyLeuValGluLysValGlnAlaAlaValGlyThrSerAlaAlaPro 311	Qy 253 GluValAlaGluValArgAlaLysLeùGluGluGlnAlaGlnGlnIleArgLeuGlnAla 272	2897 ATCTTGAGCTCCTGCCTTCGCCTTCTCTGGGTCAGTCTGCCGTTCCCTTGCTGTCTCTTG  78
QY       78 Leu       78         Db       2658 CT-GACGTAAGTGCCCACCCGACTCCCGCGCGCGCGCGCGCGC	Qy 38 TrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrp 57	Qy 1 MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20	COUNTRY: US COUNTRY: US ZIP: 77210-4433 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/819/964 FILING DATE: 28-Mar-2001 CLASSIFICATION DATA: APPLICATION NUMBER: US/09/819/964 FILING DATE: CUNKNOWN> APPLICATION NUMBER: US/09/9155 FILING DATE: CUNKNOWN> APPLICATION NUMBER: US/09/046/094 FILING DATE: OS-MAY-1997 ATTORNEY/AGENT INFORMATION: REFERENCE/DOCKET NUMBER: TAMK:177 TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 418-3000

بانوت

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Sequence 28, Application US/08726306A
Patent No. 5958684
GENERAL INFORMATION:
APPLICANT: van Leeuwen, Frederik Willem
APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
TITLE OF INVENTION: DIAGNOSIS METHOD AND REAGENTS
NUMBER OF SEQUENCES: 189
                                                                                                                                                                                                                                                                                                       CAGTGGGCCGGGCTGGAGAGGATGCAGTCGGCCGTGAGCATAAGCTCCTCCACCTCT 3736
                                                                                                                                                                                                                                                              GlnTrpAlaGlyLeuValGluLysValGlnAlaAlaValGly---ThrSerAlaAlaPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-726-306A-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/009,83
FILING DATE: 01-Jan-1996
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 96,048-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9110
TELEPAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 28:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: double TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: GB 95/20080.4 FILING DATE: 02-Oct-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/N SOFTWARE: WordPerfect 6.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 660 base pairs
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                                                                                                                                                                                                                                                                                                                                                                  ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                                                                                    LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                                                                                                                                                                                                           GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
                                                                                                                                                                                                                                                                                                                         GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
CGGCTGGGCGCGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
                                                            ACCCCGGTAGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC
                                                                          ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
                                                                                                                                                                                  CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAAGTCACCCAAGAACTGAGGGCG
                                                                                                                                                                                                                                               GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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EM: PC-DOS/MS-DOS
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RESULT 6 US-08-726-306A-28

CORRESPONDENCE

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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION UNMBER: 29,768
REFERENCE/DOCKET NUMBER: 1678
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HORNES, E. APPLICANT: UHLEN, Ma: TITLE OF INVENTION: (NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                  TOPOLOGY: lin
MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 330 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                     No.:
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               LysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeuThrProValAlaGluGlu 106
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  AAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTGACCCCGGTGGCGGAGGAG
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Matches:
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US-09-827-854-19 (1-317) x US-08-617-256-24 (1-252)
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Query Match:
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                                                                                  Alignment Scores:
                                                                                                               US-08-617-256-24
                            Best Local Similarity:
                                         Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6043031 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24, Appl Patent No. 604303
                                                                                                                                                                                                             TELEFAX: (617)227-594 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                        REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: SQ.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: DNA Diagnost
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
                                                                                                                            STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth A.
                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08,
FILING DATE: March 17,
                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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                                                                     No . :
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release "' "
                                                                                                                                                                       TYPE:
                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: March
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                                                                                                                                                                     nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kyster,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                                                                                                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.25
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96.43%
23.34%
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Patent No.
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/406,19
FILING DATE: 03-MAR-1995
ATTORNEY_AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE_DOCKET NUMBER: 3775
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 619-450-8400
TELEPAX: 619-587-5360
TELECOMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/6: FILING DATE: 18-MAR-1996 PRIOR APPLICATION DATA:
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PRIOR APPLICATION NUMBER: 08/617,256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1. CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Eh
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                             No.:
                                                                                                                                                                               TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: lin
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CITY: L
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Length:
Matches:
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Percent Similarity:
Best Local Similarity:
Query Match:
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GENERAL INFORMATION:
                                                                                       NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 2473
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEPAX: 619-587-5360
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617,227-7400
TELEPHONE: (617,227-7401
                                                              TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: K ster, H TITLE OF INVENTION: NUMBER OF SEQUENCES:
                              SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/4
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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SOFTWARE: Patent:
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/6: FILING DATE: 18-MAR-1996
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CITY: La Jolla
STRANDEDNESS:
                  TYPE:
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                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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7th Floor
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          TELEFAX: 858-587-5360
INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatibl
OPERATING SYSTEM: DOS
SOFTWARE: Patentin Rele
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SEQUENCE CHARACTERISTICS:
                                   NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002J
TELECOMMUNICATION INFORMATION:
TELEPHONE: 858-450-8400
                                                                                                                                     APPLICATION NUMBER: 08/6:
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/4(
                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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                                                                                                           FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                PRIOR APPLICATION DATA:
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| 241 GAGCGCGGCCTC 252
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                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/504, 245 FILING DATE:
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In Release #1
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                                                                                                                                                                                 08/617,256
                                                                                                                                                                                                                                                                  .0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                & McAuliffe LLP
7th Floor
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Matches:
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Indels:
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STRANDEDNESS:
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US-09-287-682-24
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Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: 1; MOLECULE TYPE: US-09-504-245-24
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                                                                                                                                                                                                                                                                                                                        ZIP: 92037-9103
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORNATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24:
TELECOMMUNICATION INFORMATION:
                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                      OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: K ster, Hubert APPLICANT: Little, Daniel P. APPLICANT: Braun, Andreas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: DNA Diagnostics Based on Mass Spectrometry NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: La Jo
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                                                                                              APPLICATION 'NUMBER: 08/406,199 FILING DATE: 03-MAR-1995
                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/287,682 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
TELECOMMUNICATION INFORMATION:

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                                                                                             ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor
CITY: La Jolla
STATE: CA
COUNTRY: USA
ZIP: 92037-9103
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
COUNTRY: MILLIANDED. MC ACTOR ATA
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24, Application US/09287679 Patent No. 6258538 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                              APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
TITLE OF INVENTION: DNA Diagno
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18 MAR-1996
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                     APPLICATION NUMBER: FILING DATE:
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96.43%
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RESULT 14
US-09-397-766-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-827-854-19 (1-317) x US-09-287-679-24 (1-252)
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Best Local Similarity:
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                             Sequence 24, Application US/09397766 Patent No. 6268144
                                                                                                                                                                                                             Patent NO. 0400...
GENERAL INFORMATION:
APPLICANT: K ster, Hubert
APPLICANT: K ster, Hubert
TITLE OF INVENTION: DNA Diagnostics
TITLE OF SECUENCES: 33
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APPLICATION NUMBER: 08/406,199
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REGISTRATION NUMBER: 34736-2002C
TELECOMMUNICATION INFORMATION:
                                                                            CITY
STATE: CA
COUNTRY: USA
COUNTRY: 92037-9103
"TP: 92037-9103
                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                 NUMBER OF SEQUENCES: 33

CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor
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TELECOMMUNICATION: INFORMATION:
TELEPHONE: (617)227-7400
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                              COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                        CITY:
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            SOFTWARE:
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TELEFAX: 619-587-5360
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RE: PatentIn Release #1.0, Version #1.25
APPLICATION DATA:
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Matches:
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Indels:
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                                                                                                                                                                        Sequence 24, Application US/0928768 Patent No. 6277573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-20021
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEPHONE: 619-587-5360
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-5941
TELEPAX: (617)227-5941
TELEPAX: (617)227-5941
                                                                                                                                          GENERAL INFORMATION: APPLICANT: K ster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
                                                          NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESSE: Heller Ehrman White
STREET: 4250 Executive Square
                                                                                                                             APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,77
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Query Match:
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FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/406,19
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 2473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (617)227-5941 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 619-450-8400 TELEFAX: 619-587-5360
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SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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              189 GluArgGlyLeu 192
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 GAGCGCGGCCTC
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                                                                   AspAlaAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla
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Command line parameters:

MODEL=frame+_p2n.model -DEV=xlp
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-DB-Published_Applications_NA -QFMT-fastap -SUFFIX-p2n.rnpb -MINMATCH=0.1
-LOOPEXT=0 -LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX-bicsum62
-TRANS-human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE-pct -THR_MX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -UTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=200000000 -USER-US09827854_[CGN_1_1_98_@trunat_11032003_101612_27578
-NCPU=6 -ICPU=3 -NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -MATIT -LONGLOG
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Maximum DB seq length: 2000000000
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    /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
    /cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
    /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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US-09-870-759-129
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Sequence 12, Appl
Sequence 129, App
Sequence 17, Appl
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US-09-827-854-12
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## ALIGNMENTS

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; ORGANISM: Homo sapiens US-09-827-854-12
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        Percent Similarity:
                                                                                                                                                                                                                      SEQ ID NO 12
LENGTH: 1156
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriak
                                                                                                                                                                                                                                                                                                                           APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing hypertriglyceridemia
FILE REFERENCE: 07180/004003
CURRENT FILING DATE: 2001-04-05
CURRENT FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                        SOFTWARE:
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100.00%
        Conservative:
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                           ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis
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; LENGTH: 1156
; TYPE: DNA
; ORGANISM: HOMO SAPI
; FEATURE:
; NAME/KEF: CDS
; LOCATION: (61)..(10
; OTHER INFORMATION:
US-09-870-759-129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Patent No. US20020177551A1
; GENERAL INFORMATION:
APPLICANT: TERMAN, David S
: TITLE OF INVENTION: COMPOSITIONS AND METHODS
: FILE REFERENCE: 870759
; CURRENT APPLICATION NUMBER: US/09/870,759
; CURRENT FILING DATE: 2002-01-14
; PRIOR APPLICATION NUMBER: US 60/208,128
; PRIOR FILING DATE: 2000-05-30
; NUMBER OF SEQ ID NOS: 166
; SOFTWARE: PATENTIN VERSION 3.1
; SEQ ID NO 129
; LENGTH: 1156
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                                                         GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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                                                                                                                             CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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; OTHER INFORMATION:
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Pred. No.:
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                                                                                                                                                                                                                                                                                                     SEQ ID
                                                                                                                                                                                                                                                                                                                    APPLICANT: Kleyn Patrick
TITLE OF INVENTION: GENES AND POLYMORPHISMS ASSOCIATED WIT
TITLE OF INVENTION: CARDIOVASCULAR DISEASE AND THEIR USE
FILE REFERENCE: 24736-2048
CURRENT APPLICATION NUMBER: US/09/802,640
CURRENT APPLICATION NUMBER: US/09/802,640
CURRENT FILING DATE: 2001-03-09
NUMBER OF SEQ ID NOS: 122
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Braun, Andreas
APPLICANT: Bonsal Aruna
                                                                                                                                                                                                                                                                                                              SOFTWARE:
                                                                                                                                                                                                                                                 ORGANISM: Homo sapien FEATURE:
NAME/KEY: CDS
LOCATION: (61)...(1014)
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CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PastSEQ for Windows Version 4.0
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US-09-827-854-8
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                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                              Sequence 8, Application US/09827854 Patent No. US20020123093A1
                                       SEQ ID NO 8
TYPE: DNA
ORGANISM: Homo
-09-827-854-8
                                                                                                                                 APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and
TITLE OF INVENTION: cholesterol
FILE REFERENCE: 07180/004003
                              LENGTH:
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RESULT 5
US-10-044-090-454
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VS-200
Patent NO. US-200
GENERAL INFORMAT
APPLICANT: Olgan
TITLE OF INVENT
FILE REFERENCE:
CURRENT FILING
CURRENT FILING
CURRENT FILING
VOUNBER OF SEQ I
SOFTWARE: PERL
SEQ ID NO 454
LENGTH: 1.291
LENGTH: 1.291
VEVE UNA
ORGANISM: HOME
FEATURE:
NAME/KEY: mis
OTHER INFORMAL
VEVE UNA
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TYPE: DNA
ORGANISM: Homo sapie
FEATURE:
NAME/KEY: misc_featu
OTHER INFORMATION: I
NAME/KEY: unsure
LOCATION: 46
OTHER INFORMATION: a
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Patent No. US20020137081A1
GENERAL INFORMATION:
APPLICANT: Olga Bandman
TITLE OF INVENTION: GENES DIFFERENTIALLY EXF
FILE REFERENCE: PA-0028 US
CURRENT APPLICATION UMBER: US/10/044,090
CURRENT FILING DATE: 2002-01-09
NUMBER OF SEQ ID NOS: 850
SOFTWARE: PERL Program
SEQ ID NO 454
LENGTH: 1291
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Best Local Similarity:
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TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Cholesterol levels with the filter perfect of 180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
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SEQ ID NO 9
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriak
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                                                                       GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: coholesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/579,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
SEQ ID NO 11
SEQ ID NO 11
LENGTH: 1156
TYPE: DATE: 156
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                                                                                                                                                                                                                                                                                                             Sequence 11, Application US/09827854 Patent No. US20020123093A1
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                    SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
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PRIOR APPLICATION NUMBER: US/60/235,720
PRIOR FILING DATE: 2000-09-27
PRIOR PELICATION NUMBER: US/60/235,840
PRIOR APPLICATION NUMBER: US/60/235,863
PRIOR APPLICATION NUMBER: US/60/235,863
PRIOR EILING DATE: 2000-09-27
NUMBER OF SEQ ID NO5: 2276
SOPTWARE: Patentin version 3.0
SEQ ID NO 760
LENGTH: 1157
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CURRENT FILING DATE: 2001-09-18
PRIOR APPLICATION NUMBER: US/60/233,617
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: US/60/234,052
PRIOR APPLICATION NUMBER: US/60/234,953
PRIOR APPLICATION NUMBER: US/60/234,923
PRIOR FILING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US/60/235,134
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,638
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APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
FILE REFERENCE: 44921-5028-W0
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2244
LENGTH: 1157
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
COTHER INFORMATION: Genbank Accession No. US
US-09-880-107-2244
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing hyp
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILLING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEO ID NOS: 19
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Best Local Similarity:
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ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
APPLICATION NUMBER: US/09/827,854
CURRENT APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-104-06
PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 10
LENGTH: 1156
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Sequence 5, Application US/09967013

Patent No. US20020045840A1

GENERAL INFORMATION: Jr. Vincent p. APPLICANT: Stanton, Jr. Vincent p. TITLE OF INVENTION: METHOD FOR GENETIC ANALYSIS OF APOE DNA TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: 11926-022001

CURRENT APPLICATION NUMBER: US/09/967,013

CURRENT APPLICATION NUMBER: 60/206 613

PRIOR APPLICATION NUMBER: 60/206 613

PRIOR FILING DATE: 2000-05-23

NUMBER OF SEQ ID NOS: 91

SOFTWARE: FrastsEQ for Windows Version 4.0

SEQ ID NO 5

LENGTH: 41907

TYPE: DNA

ORGANISM: Homo sapiens
                    Alignment Scores: Pred. No.:
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Length:
Matches:
Conservative:
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	aGlnalaArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrAr 	uGlnLeuThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaA 	-ArgAlaLeumetaspGluThrmetLysGluLeuLysAlaTyrLysSerGluLeuGl 	CCCCATCCCAGCCCTCTCCCCGCCTCCCACTGTGCGACACCCTCCCGCCCTCTCGGC		GCCTCTGCCCCGTTCCTCTCTCCCCTCTTGGGTCTCTGGCTCATCCCCCATCTCGCC		PCGTCTCTGCCTCTGCCTCTGCATCTGCTCTCTGCATCTGTCTCTGTCTCTCTTCTC		CCTCCCAAAGTGCTGGGATTAGAGGCATGAGCCACCTTGCCCGGCCTCCTAGCTCCTT		CTCACTGTGTTGCCCAGGCTGGTCTTGAACTTCTGGGCTCAAGCGATCCTCCCGCCTC		CGTCCTGGCTCTGTCTCTGTCCCTTCCCTAGCTCTTTTATATAGAGACAGAGAGAG		TGGAATTCTCTCTCAGCTTTGTCTCTCTCTCTTCCTTTGAGTCAGTC		CCTGGGTCTCTGCTGGTTCTAGCTTCCTCTTCCCATTTCTGACTCCTGGCTTTAGCTCTC		GGCGGCTATACCTCCCCAGGTCCAGGTTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGGG						LeuargGlnGlnThrGluTrpGlnSerGlyGlnArgTrpGluLeualaLeuGlyArgPhe	PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProG 	-854-19 (1-317) x US-09-967-013-5 (1-41907)	al Similarity: 59.80% Mismatches: 6 86.15% Indels: 194 10 Gaps: 1
	gG1 138      CGG 21275	Al 118    GC 21215	luG1 98       aGGA 21155	CG 21095	78	CG 21035	78	TCG 20975	78	CT 20915	78	GG 20855	78	GT 20795	<sup>'</sup> 78	CT 20735	78	TC 20675	78	GG 20615	78	TG 2	78	N	er 71	he 51    TT 20435	lu 31     AG 20375		

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Percent Similarity:
Best Local Similarit.
Query Match:
DB:
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CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1999-03-12
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 133
LENGTH: 786
TYPE: DNA
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic
FILE REFERENCE: PA104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 133, Application US/09925302 Patent No. US20020044941A1
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                                                                                                                                                            Similarity:
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867.50
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                                  US-09-827-854-19 (1-317) x US-09-964-824A-374 (1-478)
                                                                     Best Local Similarity:
Query Match:
                                                                                            Percent Similarity:
                                                                                                                                                     ; NAME/KEY: misc_feature
; LOCATION: (1)...(478)
; OTHER INFORMATION: n=a
US-09-964-824A-374
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US-09-964-824A-374/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 374, Application US/09964824A Patent No. US20020102531A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/964,824A CURRENT FILING DATE: 2001-09-27
                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Cancer TITLE OF INVENTION: Sets FILE REFERENCE: 689290-73
                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US/60/236,028 PRIOR FILING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Horrigan,
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                                                                                                                                                                                                    ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                      LENGTH: 478
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96.92%
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Query Match:
DB:
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NAME/KEY: unsure
LOCATION: (1)...(478)
OTHER INFORMATION: n = a or c or g or
US-09-880-107-2491
                                                    В
                                                                                                                                                                                  Score
                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
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US-09-880-107-2491/c
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                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 2491
LENGTH: 478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Horne,
                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                              . No.:
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Search completed: March 15, 2003, 03:05:08 Job time : 80.8753 secs

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MODEL=frame+_D2n.model -DEV=xlp

-Q-cgn2_1/USPT0_Spool_VS09827854/runat_11032003_101610_27495/app_query.fasta_1.3576
-DB=EST -QFMT=fastap -SUFFIX=p2n.rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXI=0

-UNITS=blits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFMT=pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXIEN=200000000

-USER=US09827854 @CGN_1_1_8825_@runat_11032003_101610_27495 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAXIT -LONGLOG -DEV_TIMEOUT=120
-WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

## ALIGNMENTS

AUTHORS TITLE	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BI670367	RESULT 1
NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 1027)	Homo sapiens	human.	EST.	BI670367.1 GI:15584600	BI670367	mRNA sequence.	603292738F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5312024 5',	BI670367 1027 bp mRNA linear EST 12-SEP-2001		

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Email: cgapbs r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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Contact: Robert St
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/db_xref="taxon:9606"
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/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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BG763371
BG763371.1 GI:
EST.
                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (
                                                                                                                    Plate: LLCM1718 row: b column: High quality sequence stop: 817.
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National Institutes of Health, Mammalian
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 842)
                                                                                                                                                             DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                 Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                              Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                 human.
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                                                                                                                                                 nttp://image.llnl.gov
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4860585"
/clone_lib="NIH_MGC_49"
/tissue_type="melanotic melanoma, high
/lab_host="DH10B (phage-resistant)"
                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                             GI:14074024
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ECCRI; CDNA made by Ollgo-dT priming. Directionally cloned
into ECCRI/XhoI sites using the following 5' adaptor:
GGCACGAG(G). Size-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
ILRT (Life Technologies). Note: this is a NIH_MGC
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TITLE
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                                                                           GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                              ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCCAGGATGCCCAGGCCAAGGTG
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Location/Qualifiers
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
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/clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                                                                                                                                                       mRNA sequence.
BG761746
BG761746.1 GI:
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTP/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium ()
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 938)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collections
                                                               Unpublished (1999)
Contact: Robert St
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602717942F1 NIH_MGC_49 Homo s
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Best Local Similarity:
GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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                                                                    AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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quality sequence stop:
Location/Qualifiers
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/clone_Iib="NIH_MGC_49"
//tissue_type="melanotic melanoma, high MDR (cell line)"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/lab_host="Organ: skin; Vector: pOTB7; Site_1: xhoT; Site_2:
/note=""Organ: skin; Vector: pOTB7; Site_1: xhoT; Site_2:
/note="Contain the pot of the following 5; adaptor: into EcoRI/XhoI sites using the following 5; adaptor:
//organ: size-selected >500bp for average insert size
/labb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-CDNA synthesis kit (Stratagene) and Superscript
II RT (Life Technologies). Note: this is a NIH_MGC
Library. | "
71 a 272 c 374 g 121 t
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                              DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11762 row: a column: 11
High quality sequence stop: 762.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Strayaki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 942)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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603249241F1 NIH_MGC_96
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                                                                                 /note-"organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site-1: BamHI; Site_2: SalI-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length; clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library." a 284 c 361 g 120 t
                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5301010"
                                                                                                                                                                                                                     /clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
  3.39e-117
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603615713F1 NIH
mRNA sequence.
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BM042094.1 GI:
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KEYWORDS
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (
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/lab_host="pHI()B (phage-resistant)"
/note="organ: skin; Vector: pOTBF; Site_1: XhoI; Site_2:
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1 (bases 1 to 927)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov

Tissue Procurement: ATCC
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Contact: Robert Strausberg, Ph.D.
Email: Cgapbs-remail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), S
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                  LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                  CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCAGGTCACCCAGGAACTGAGGGCG
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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/db_xref="taxon:9606"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                      CTGATGGACGAGCCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
                                                                                                                                                     Unpublished (1999)
Contact: Robert Strausberg, FEmail: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTP/DTP
cDNA Library Preparation: Li
                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I Mammalla; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 817)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
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Plate: LLCM1612 row:
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                                                                                                            cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information of through the I.M.A.G.E. Consortium/LLNL at:
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/Organism="Homo sapiens"
/Ob_xref="taxon:9606"
/Clone="IMAGE:4760827"
/clone_lib="NIH_MCC_40"
/tissue_type="carcinoma, cell line"
/tasue_type="carcinoma, cell line"
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                                                                       LeuGlnGluArgAlaGlnAla--TrpGlyGluArgLeuArgAlaArgMetGluGluMetG
                                                                                                               CCCCTGGTGGAACAGGGCCGCGTGCGGGCCTGCACTGTGGGCTCCCTGGCCGGCAG-CCG
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CTGTCTGAGCAGGAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI), i
Toshiyuki and Piero Carninci (RIKN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11726 row: g column: 10
High quality sequence stop: 783.
Location/Qualifiers
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Contact: Robert Strausberg, Ph.D
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NIH-MGC http://mgc.nci.nih.gov/
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC
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National Institutes of Health, Mammalian Gene Collection (MGC)
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            SerargThr-ArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
                                                                     LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
                                                                                                                               ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
                                                                                                                                                                         CAGGCCGGGGCCCGAGGGCCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
                                                                                                                                                                                       GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                                                                                                                                                                                                                                 LysLeuArgGlnArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                                                                                                                                                                                                                                                      GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHísLeuArg
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                                                                                                                CCCCTGGTGGAACAGGGCCGCGTGCGGGCCGC - ACTGTGGGCTCCCTTGGCGGCAG - CCG
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Directionally cloned into ECORI/Xho! sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZaP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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1171.50
96.23%
95.09%
73.91%

    CGCCTGGACGAGGTGAAGGAGCAGGTGCGCGCCAA

                                                        -GGGCGAGCGGTGCGCGCGCGGATGGAGGAGATGGGC
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.

Email: cgapbs r@mail.nih.gov

Tissue Procurement: Miklos Pakkovits, M.D., Ph.D.

CDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh

Toshiyuki and Piero Carninci (RIKEN)

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can

found through the I.M.A.G.E. Consortium/LLNL at:

http://lmage.llni.gov

plate: LLAM10672 row: e column: 21
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National Institutes of Health, M
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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//db_xref="Laxon:9606"
//clone="IMAGE:4793084"
//clone=lib="NIH_MGC_96"
//tissue_type="hypothalamus"
//tissue_type="hypothalamus"
//lab_host="hyll08"
//lab_host="hyll08"
//lab_host="brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag pBluescript ROT 5. This is a primary library enriched normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

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                                                                                                                                          Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih!gov

Tissue procurement; Miklos Palkovits, M.D., Ph.D.

cDNA Library Preparation: Michael J. Brownstein (NHGRI), incomparation: Michael J. Brownstein (LLNL)

CDNA Library Arrayed by The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC [clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov

Plate: LLAM11691 row: 1 column: 04
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National Institutes of Health, Mammalian
Unpublished (1999)
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BI551475.1 GI:15438787
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                               quality sequence stop:
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:52/4003"
/clone_lib="NIH_MGC_95"
/tissue_type="hippocampus"
/lab_host="DH10B"
                                                                                                                  Location/Qualifiers
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| Homo sapiens cDNA clone IMAGE:5274003 5',
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Tissue Procurement: DCTD/DTP
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National Institutes of Health, M
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/lab_host="hHl0B (phage-resistant)"
/note="Organ: skin: Vector: poTBP; Site_1: XhoI; Site_2:
/note="Organ: skin: Vector: poTBP; Site_1: XhoI; Site_2:
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/db_xref="taxon:9606"
/clone="IMAGE:5557004"
/clone_lib="NIH_MGC_112"
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                                                                                                                                                                                             Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                         Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BG706129 907 bp r
602669093F1 NIH_MGC_96 Homo sapiens
                                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov plate: LLAM10669 row: i column: 23 High quality sequence stop: 832.
                                                                                                                                                                                                                                                                                                                                    NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                    Mammalia; Eutheria;
1 (bases 1 to 907)
                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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BG706129.1 GI:13981169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence
                                                                                                                     quality sequence stop: 0
            /organism-"Homo sapiens"
/db_xref-"taxon:9606"
/clone="IMAGE:4792030"
/clone_lib-"NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host-"DH10B"
                                                                                                        .907
 brain;
Vector:
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pBluescriptR (modified
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Brownstein
                                                                                                                                                                                                                                                                                                                                       Gene
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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ORIGIN
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                                  lGlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuAr
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GCAG-CGGACCCGG---ACGCTGGACGAGTG-AAGGAGCAGTGGGGAAGTGTGCGC
                                                                                                                                                                                    GCAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCG
                                                                                                                                                                                                                        CCGGCTGGGCGGGCATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGT
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a 260 c 358 g 117 t 1 others
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1125.50
94.628
93.468
71.018
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Matches:
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Search completed: March 14, Job time: 1293.69 secs 2003, 20:14:18

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Command line parameters:

-MODEL-frame+_p2n.model -DEV=xlp
-Q-cgn2_1/USPTO_spool/US09827854/runat_11032003_101609_27476/app_query.fasta_1.3576
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

5 100.0 954 24 AAD26035 5 100.0 1110 7 AAN60409 5 100.0 1147 22 AAP84315 5 100.0 1156 24 AAD22048 5 100.0 1156 24 AAD22048 7 99.5 1156 24 AAD22049 7 99.5 1156 24 AAD22049 7 99.5 1156 24 AAD22049 7 99.5 1156 24 AAD22049 7 99.5 1157 24 ABD2574 6 99.4 1157 24 ABD2574 6 99.4 1157 24 ABD2574 6 99.4 1157 24 ABD2574 6 99.4 1157 24 ABD2574 6 99.4 1157 24 ABD25045 6 99.4 1156 22 AAP84316 7 99.1 1156 22 AAP84316 7 99.2 1156 24 AAD22047 7 99.1 1156 24 AAD22047 7 99.1 1156 24 AAD22047 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22050 7 99.1 1156 24 AAD22075 7 99.3 1107 19 AAX75264 7 98.1 1076 24 AAD22075 7 99.3 1107 19 AAX752081 7 87.4 1076 24 AAD22075 8 71.4 1076 24 AAD32075 8 71.4 1076 24 AAD32075 8 71.5 126 19 AAX718068 8 71.6 10 AAX718068 8 71.6 10 AAX718068 8 71.7 786 21 AAX718068 8 71.8 1076 24 ABD3599 8 42.8 600 20 AAX99599 8 42.8 600 20 AAX99599 9 39.4 478 24 ABD35049 9 39.4 478 24 ABD35049 9 39.5 409 22 AASS3049	Score Match De ID    Score Match Length DB   ID   D
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## ALIGNMENTS

RESULT 1 AAD26035

AAD26035 standard; cDNA; 954 BP

AAD26035;

26-MAR-2002 (first entry)

Human apolipoprotein E (APOE) cDNA.

Human; antilipaemic; neuroprotective; nootropic; genetic variant; APOE; apollpoprotein E; haplotyping; familial dysbetalipoproteinaemia; therap; genotyping; type III hyperlipoproteinaemia; Alzheimer's disease; SNP; atherosclerosis; single nucleotide polymorphism; chromosome 19q13.2; ss. therapy;

Homo sapiens

Key

Location/Qualifiers

US-09-827-854-15 (1-317) x AAD26035

(1-954)

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Best Local
Query Match
DB:
                                                                                   Score:
                                 Percent Similarity:
                                                                                                   The patent discloses novel genetic variants of human apolipoprotein E (APOE) gene. The invention also relates to compositions and methods for haplotyping and/or genotyping the APOE gene. The haplotyping comethods of the invention are useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with APOE activity, e.g. catherosclerosis, and Alzheimer's disease. They are useful to validate APOE as a candidate agent for treating a specific condition or disease predicted to be associated with APOE activity. Genotyping or disease predicted to be associated with APOE activity. Genotyping or disease predicted to be associated with APOE activity. Genotyping or disease predicted to be associated with APOE activity. Genotyping or disease predicted to be associated with APOE activity. The present sequence is a cDNA encoding human APOE protein.
                                                                                                                                                                                                                                                                                                           Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotion pair at specific polymorphic sites for two copies of gene -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDS
                                                                                   Sequence
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                      Similarity:
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Ovarian tumour marker gene; human; overexpression; upregulation; epithelial tumour; cancer; diagnosis; prognosis; disease monitoring; identification; serous cystadenoma; broderline serous tumour; serous cystadenocarcinoma; mucinous cystadenocarcinoma; mucinous cystadenocarcinoma; mucinous cystadenoma; borderline mucinous tumour; endometrioid carcinoma undifferentiated carcinoma; clear cell adenocarcinoma; cystadenofibroma; adenofibroma; Brenner tumour; serial analysis of gene expression; SAGE; immune response pathway; cell proliferation regulation; protein folding;
                                                  mmune response pathway; cell
membrane localised; secreted;
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                                                                                                                                                                                                                                   Apolipoprotein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detecting an ovarian tumour in a patient, for identifying an individual CC at increased risk for developing ovarian cancer, in prognostic tests for CC assessing the relative severity of ovarian cancer, in tests for CC monitoring a patient in remission from ovarian cancer and in tests for CC monitoring a patient in remission from ovarian cancer and in tests for CC minitoring disease status in a patient being treated for ovarian cancer. CC The methods can additionally be used to identify a particular tumour as being an ovarian tumour (i.e., an epithelial ovarian tumour selected from CC serous cystadenoma, borderline serous tumour, serous cystadenocarcinoma, condentified cystadenocarcinoma, cystadenosticnoma, cucinous cystadenocarcinoma, cystadenofibroma, adenofibroma and Brenner CC tumour. The ovarian tumour marker genes of the invention were identified cusing SAGE (serial analysis of gene expression) and were found to be coverapressed in a broad variety of ovarian epithelial tumour cells to cell implicated in immune response pathways, in the regulation of cell coalised or secreted. In addition to their use as diagnostic and CC proteins may be used as therapeutic targets for the treatment and CC proteins may be used as therapeutic targets for the treatment and CC proteins may be used as therapeutic targets for the treatment and CC proteins may be used as therapeutic targets for the treatment and CC proteins may be used as therapeutic targets for the treatment and CC proteins may be used as therapeutic targets for the treatment and CC proteins may be used as the coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour may be used as the coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour marker genes or their encoded coarian tumour marker
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                                                               LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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or preventing
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05-APR-2001;
The present sequence is a human apolipoprotein E (apoE) isoprofein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atheroscierosis, treating or regressing atherosclerosis without inducing hypertriglycerideemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (I receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or h
                                                                                                                                                              WPI; 20
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05-APR-2001;
                                                                                          The present sequence is a human apolipoprotein E (apoE) allele, apoE2** DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (L receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or hadden.
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                                                                                                                                                                                                                                                                                                                                                         Sequence 1156
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE2 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (Preceptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New apolipoprotein E polypeptide and nucleic acid, cholesterol, delaying the onset of or treating athmammal, without inducing hypertriglyceridemia -
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PT New apolipoprotein E polypeptide and nucleic acid, useful for lowering PT cholesterol, delaying the onset of or treating atherosclerosis in PT mammal, without inducing hypertriglyceridemia -
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Claim 14; Page 82; 91pp; English.

The present sequence is a human apolipoprotein E (apoE) allele, apoE2\* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing the present apolity and the sequence of the protein and protein an amamal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LDL) receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or having a defect in remnant removal.

Sequence 1156 BP; 208 A; 367 C; 432 G; 149 T; 0 other;

Pred. No.: 5.3e-114 Length: 1156
Score: 1577.00 Matches: 316
Percent Similarity: 99.68% Conservative: 0
Best Local Similarity: 99.68% Mismatches: 1
Query Match: 99.50% Indels: 0
DB: 04.500 Gaps: 0

US-09-827-854-15 (1-317) x AAD22051 (1-1156)

240	LeuGInGluArgAlaGInAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly:	221	Qу
720		661	DЬ
220	ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro	201	Qy
660		601	Db
200	GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly	181	Qy
600		541	Db
180	LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr	161	Qy
540		481	ρb
160	GlnAlaMetLeuGlyGlnSerThrGluGluIeuArgValArgLeuAlaSerHisLeuArg	141	Qy
480	CGCCTGGGCCCGACATGGAGGACGTGTGCGGCCCTGCTGCAGTACCGCGGCGAGGTG	421	рь
140		121	Qy
420	ACCCCGGTGGCGGAGGAGGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGCAGGCGCAGGCC	361	Db
120		101	Qy
360		. 301	Db
100	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu	81	Qy
300		241	ДD
80	LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla	61	Qy
240		181	Db
60	GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr	41	Qy
180	GAGCAGAGCGGTGGAGACAGAGCCCGAGCCGGAGCCGAGCAGACCGAGTGGCAGAGC	121	Db
40	GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer	21	Qy
120	ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG	61	Db
20	MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal	1	γo

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RESULT 10
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Recombinant human apolipoprotein-E (ApoE) (AAR86791) can be produced by preparing a genetic construct (contg. an ApoE-encoding sequence, e.g. AAT06957, and flanking regulatory sequences enabling the protein to be expressed in insect cells), which is then introduced into a Manduca sexta larva (using a recombinant Autographica californica nuclear polyhedrosis virus) and recovering the protein from the haemolymph of the larval host. The ApoE produced is in a form sufficiently complexed with lipids to be biologically active, which cannot be achieved in insect cell cultures, and can therefore be used in therpeutic applications.
                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                Manduca
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Manduca sexta; Autographica californica nuclear
haemolymph; lipid complex; biologically active;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
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                                                                                                                                                                                                                                                                                                     of recombinant apo:lipoprotein E in insects sexta larvae with recombinant Autographica polyhedrosis baculovirus vector.
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    by infecting californica

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RESULT 11
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                                                                                                                                                                                                                                                                                                                      expression of two or more genes represented in ABN93503-ABN97455 in a tissue sample. The method of the invention has hepatotropic, and cytostatic activity. The method is useful for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma and metastatic liver carcinoma in a patient. The method is useful for identifying expression profiles which serve as useful diagnostic markers as well as markers that can be used to monitor disease states, disease progression, drug toxicity, drug efficacy and drug metabolism.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                        Sequence 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates progression of liver of
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                                                                                                                                                                                                                                                                                                                                                                                            CC progression of benign prostatic hyperplasia (BPH), or screening (II) for cor identifying an agent that modulates the onset or progression of BPH. CC The method is based on changes in gene expression in BPH tissue isolated from patients exhibiting different clinical states of prostate cC hyperplasia as compared to normal prostate tissue. (I) comprises CC hyperplasia as compared to normal prostate tissue. (I) comprises CC trom the subject that are differentially regulated compared to normal prostate cells. (II) comprises preparing a first gene expression profile CC prostate cells or BPH-like cell population, exposing the cells to the agent, preparing a second gene expression profile of the agent exposed cells, and comparing the first and second gene expression profiles. (I) is useful for diagnosing the onset or progression of BPH. (II) is useful for dentifying an agent that modulates the onset or progression compared to the expression of gene in the specification in the tissue or cells to the expression level in a tissue or cells, by comparing the expression clevel of genes given in the specification in the tissue or cells to the capression levels of at least one gene in the tissue or cell sample compared to the expression level in a tissue or cell sample compared to the expression level in BPH. Agents using (II) are useful for treating BPH or prostate cancer. ABK64106-ABK64860 represent human contact the capter of the invention.
                                                                                                                                                                                                                                                                   Y Match:
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30-MAY-2001; 2001WO-US10838

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cc agent to be tested for anti-neoplastic activity, determining a change in comprises a sequence (S) selected from 8447 sequences (given bed) as indentical to (S), where a change in comprises a sequence (S) selected from 8447 sequences (given in ABL61664 to ABL70110), or is at least 95% identical to (S), where a change in comprises a sequence (s) selected from 8447 sequences (given in ABL61664 to ABL70110), or is at least 95% identical to (S), where a change in compression is indicative of anti-neoplastic activity. (I) has cytostatic cativity and can be used in gene therapy. MI can be used for screening an anti-neoplastic agent, and can be used for producing a product which cis the data collected with respect to the anti-neoplastic agent as a cresult of MI, and the data is sufficient to convey the chemical concervature and/or properties of the agent. MI can be used in the tracture and/or properties of the agent. MI can be used in the creating accordance accordance accordance accordance accordance. Cc adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer, infiltrating ductal cancer, carcinoma, papillary carcinoma and Wilm's tumour.
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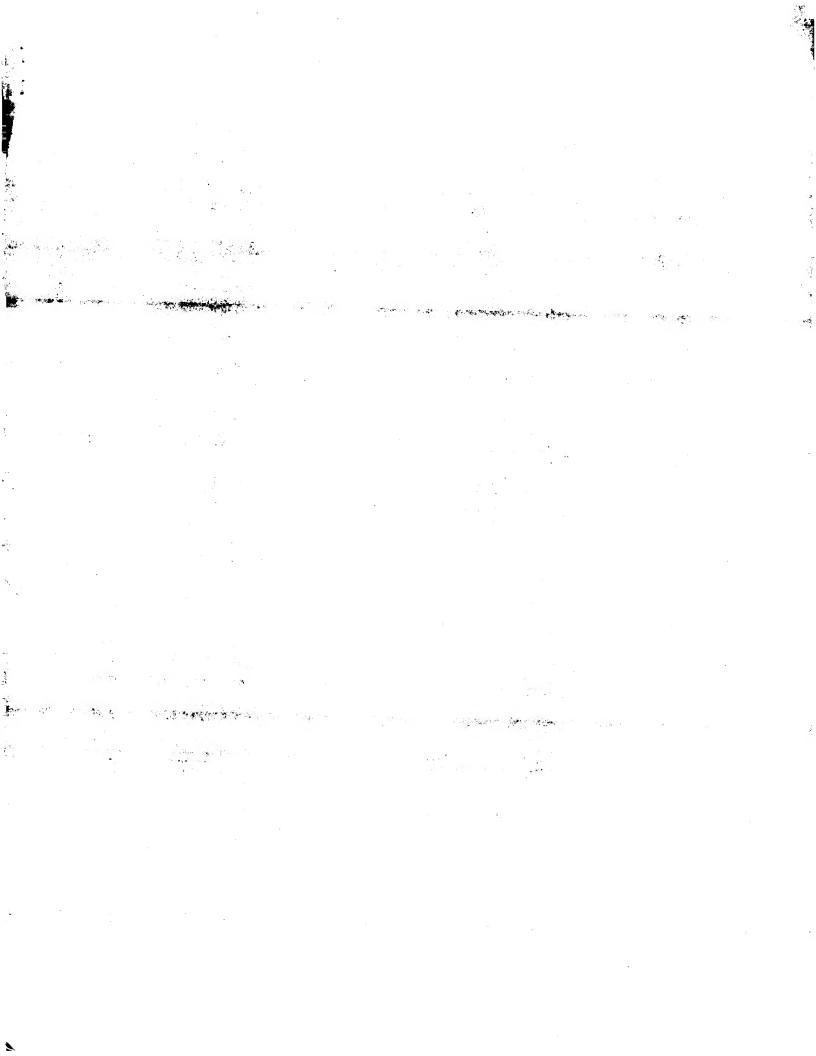
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PMetG       CATGG	ArgLeud             CGGCTGC	Qy 121 Db 421
RGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla 1  -	ThrProVal	Qy 101 Db 361
ThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluGlnLeu	LeuMetAsp	Qy 81 Db 301
NVAlGInGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 8 	LeuSerGlu(	Qy 6 Db 24
GluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 6 	GlyGlnArgTr               GGCCAGCGCTG	Qy 41 Db 181
GluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 4 	ര – ഒ	Qy 21 Db 121
1TrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20 	MetLys           ATGAAG	Oy 6
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1.09e-113 Length: 1156 1573.00 Matches: 316 99.68% Conservative: 0 99.68% Mismatches: 1 99.24% Indels: 0 22 Gaps: 0	Scores: milarity: Similarity h:	Alignment Pred. No.: Score: Percent Si Best Local Query Matc DB:
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ition relates to an ApoE humanised mammalian cell. The is the coding sequence for human ApoE4, which was us the present invention. The ApoE humanised mammalian screening for agents useful for treating or preventings and arteriosclerosis.	resent invent sequence method of used for imer's dise	CC The process of the control of the
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nd mammalian cell useful for screening for agents or preventing Alzheimer's disease and	poE humaniz l for treat iosclerosis	New use art
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Command line parameters:

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-LIST=45 -DCCALIGN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
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-USER-US09827854_@CGN_1_1_224_@runat_11032003_101611_27510 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEDUT=120
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/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
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US-09-849-389-6
US-08-617-256-24
US-09-287-141-24
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US-07-709-949-1
; Sequence 1, Ap
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Patent NO. 34,2007; GENERAL INFORMATION:
; APPLICANT: Attie, Alan D
; APPLICANT: Gretch, Daniel G
; APPLICANT: Sturley, Stephen L
; APPLICANT: Beckage, Nancy E
; APPLICANT: Production of Re
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                                                                  COUNTRY: USA
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/709,949
FILING DATE: 1991064
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 9629691801
   TELEFAX: (608);
INFORMATION FOR SEQ
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                                   TELEPHONE: (608) 251-5000
                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: PROTITLE OF INVENTION: In:
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Qua:
STREET: P.O. BO
CITY: Madison
STATE: WI
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SEQUENCE CHARACTERISTICS: LENGTH: 1157 base pairs

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AUTHORS: McLean, J
JOURNAL: Journal o
VOLUME: 259
PAGES: 6498-6504
DATE: 1984
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Best Local Similarity:
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LOCATION INFORMATION:

MCLean, J W
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ORIGINAL SOURCE:
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LOCATION: 62..1015
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                                                                                                                                                                                                                                                                                                                               GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
                                  LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                            GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                          LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                                CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
                                                                                                                                                                                                            LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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                       AAGCTGCGTAAGCGGCTCCTCCGCGATCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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Journal of Biological Chemistry
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                                                                                                                  COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Flopy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: BATENTIN Release #1.0, Ver

CORRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/949,155

FILING DATE: CONCURRENTLY Herewith

CLASSIFICATION DATA:

APPLICATION NUMBER: US 60/027,338

FILING DATE: 11-OCT-1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/046,094

FILING DATE: 09-MAY-1997

APTORNEY/AGENT INFORMATION:

NAME: Hibler, David W.

REGISTRATION NUMBER: 41,071

REFERENCE/DOCKET NUMBER: TAMK:177
                                     TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1126 base pairs
                                                                                                                                                                                                                                                                                                                                                                STATE: LA
COUNTRY: US
77210-4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Piedrahita, Jorge A
APPLICANT: Bazer, Fuller W
TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: Generation of Transgenic
NUMBER OF SEQUENCES: 51
                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 5: CORRESPONDENCE ADDRESS:
TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: lin
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CITY: H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity:
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                                                                                                     HisLeuArgLysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeu 177
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GlyGlnProLeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGlu 237
                                                                                                                                                                                                                           ArgLeuGlyProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAla
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                                                  CGCACCAAAGTGGAGGAGCAGGGCAGCCAGTTGCGCCTGCAGGCCGAGGGGATTCCACGCC
                                                                 ArgAlaLysLeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAla
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US-09-819-964-5
; Sequence 5, Application
; Patent No. 6369294
; GENERAL INFORMATION:
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TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                   . No.:
 111
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/949,155
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/046,094 FILING DATE: 09-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Compositions and Methods for the Generation of Transgenic Animal Species
                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ
                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Piedrahita, Jo
Bazer, Fuller
                           GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu 37
                                                                                 NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TA
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/819,964 FILING DATE: 28-Mar-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
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STATE: TX
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                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
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RESULT 4
US-08-949-155-51
                                                                                                                                             Sequence 51, Applicat Patent No. 6271436 GENERAL INFORMATION:
                                                        APPLICANT: Piedrahita, Jorge A
APPLICANT: Bazer, Fuller W
TITLE OF INVENTION: Composition
TITLE OF INVENTION: Generation
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE AND
STATE: TX
COUNTRY: US
ZIP: 77210-4433
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                                      Houston
                                                                                                                                                                     Application US/08949155
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Generation of Transgenic
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Best Local Similarity:
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FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 51
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
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PRIOR APPLICATION DATA:
US 60/027,338
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APPLICATION NUMBER: US/08/949,155
FILING DATE: Concurrently Herewiti
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
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TOPOLOGY: lir
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                           CCCTCCTGGCGAACCGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
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Matches:
Conservative:
Mismatches:
Indels:
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US-09-819-964-51
Sequence 51, Application US/09819964
Patent No. 6360294
GENERAL INFORMATION:
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                                                               ARPLICANT: Piedrahita, Jorge A
APPLICANT: Piedrahita, Jorge A
Bazer, Fuller W|
Bazer, Fuller W|
TITLE OF INVENTION: Compositions
Generation of
               CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE
STREET: P.O. Box 4433
                                                            NUMBER OF SEQUENCES: 51
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CITY: Houston
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Best Local Similarity:
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SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
TYPE: nucleic acid
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FILING DATE: <Unknown>
APPLICATION NUMBER: US 60/046,09
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: HAbler, David W.
REGISTRATION NUMBER: 41,071
                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
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MEDIUM TYPE: Floppy disk
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FILING DATE: 28-Mar-2001
CLASSIETCATION: CURKNOWN>
R APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
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Sequence 28, Application US/08726306A
Patent No. 5958684
GENERAL INFORMATION:
APPLICANT: van Leeuwen, Frederik Willem
APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
                APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
TITLE OF INVENTION: DIAGNOSIS METHOD AND REAGENTS
NUMBER OF SEQUENCES: 189
                                                                                                                                                                                                                                                                                   GlnTrpAlaGlyLeuValGluLysValGlnAlaAlaValGly---ThrSerAlaAlaPro
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Best Local Similarity:
Query Match:
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REFERENCE/DOCKET NUMBER: 96,048-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 28:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: GB 95/20080.4
FILING DATE: 02-0ct-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
FILING DATE: 01-Jan-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatib
OPERATING SYSTEM: PC-DOS/
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
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STREET: 1 Financial Center
CITY: Boston
STATE: MA
COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 660 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                                                                                                                                                                                 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                                                                                                                                                                                                                                                                                              GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                                 ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120
                                                                                                                                                                                                CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCAAGTCACCCAAGAACTGAGGGCG
                                                                                                                                                                                                                                                                 GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                                                                                                                                                                                                                                                                                                  GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
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                                                                 ACCCCGGTAGCGGAGGAGCGCGGGCACGGCTGTCCAAGGAGCTGCAGACGGCGCAGGCC
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                                              US-09-827-854-15 (1-317) x US-07-849 389-6 (1-330)
                                                                        Query Match:
DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION UNMER: 29,768
REFERENCE/DOCKET NUMBER: 1678
TELEPHONE: (703)836-9300
TELEPHONE: (703)836-9300
TELEFAX: (703)633-4109
                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                              Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: HOKNES, APPLICANT: UHLEN, Mathias
                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 330 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/849,389
FILING DATE: 19920519
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                              MOLECULE TYPE:
ORIGINAL SOURCE:
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MEDIUM TYPE: Floppy disk
                                                                                                                                 No.:
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OPERATING SYSTEM:
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                                                                                                                                                                                            NAME/KEY:
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LysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeuThrProValAlaGluGlu 106
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SYSTEM: PC-DOS/MS-DOS
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            Query Match:
                        Best Local Similarity:
                                   Percent Similarity:
                                                 Score:
                                                               Pred. No.:
                                                                           Alignment Scores:
                                                                                                     US-08-617-256-24
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                                                                                                                                                                                          TELEFAX: (617)227-594
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406
FILING DATE: March 17, 199
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth A.
REGISTRATION NUMBER: 35,43
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                   TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kvster, Hubert TITLE OF INVENTION: DNA D
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CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61
                                                                                                                             TOPOLOGY:
                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US, FILING DATE: March 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                         (617)227-5941
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18, 1996
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Indels:
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US-09-827-854-15 (1-317) x US-08-617-256-24 (1-252)

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Alignment Scores: Pred. No.:
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ARE DATE: 03-MAR 1995
ATTORNEY/AGENT INFORMATION:

NAME: Seidman, Stephanie L

REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002D
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEPHONE: 619-587-5360
TELECANX: 619-587-7360
TELECHAN: 617)227-7400
TELEPHONE: (617)227-7941
TELEPHONE: (617)227-5941
TELEFAX: 617)227-5941
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Patent No. 6197498
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                                                                                                                                                                                                                                                                                                                         TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/61
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/41
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CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/287,141
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                          STRANDEDNESS:
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SOFTWARE: Patentl
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Best Local Similarity:
Query Match:
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Patent No. 6221601
                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617
FILING DATE: 18 -MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406
FILING DATE: 03 -MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                        TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 2:
                         SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
                                                                                                                                                   TELEPHONE: 619-450-8400
                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
ADDITION NUMBER: US/09/431,613
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: K ster, Hubert
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CITY: La
STATE: CA
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OPERATING SYSTEM: DOS
OPETWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 GAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGCAAGCTGCGTAAGCGGCTCCTCCGC 180
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TELEFAX: 619-587-5360
                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 24
                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 92037-9103
              STRANDEDNESS:
                                                                                                                                                                                                               NAME:
TOPOLOGY:
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7th Floor
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Indels:
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: US-09-431-613-24
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE,DOCKET NUMBER: 24736-2002J
TELECOMMUNICATION INFORMATION:
TELEPHONE: 858-450-8400
TELEPAX: 858-587-5360
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                 OPERATING SYSTEM: DOS
SOFTWARE: Patentin Release #3
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/09/50
FILING DATE:
CLASSIFICATION:
                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/61
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
                                                                                                      APPLICATION NUMBER: 08/4
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
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                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
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                                                                                                                                                                                                                                                                                                                                                CITY: La
STATE: CA
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92037-9103
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7th Floor
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Matches:
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; STRANDEDNESS:
; TOPOLOGY: line
; MOLECULE TYPE: C
US-09-504-245-24
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Best Local Similarity:
Query Match:
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US-09-287-682-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                            FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
FILING DATE: 03-MAR-1995
ATTORNEY/ACENT INFORMATION:
NAME: Seidman, Stephanie L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
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APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
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                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                       TLITLE OF INVENTION: DNA Diagnostics Based on NUMBER OF SEQUENCES: 33
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REFERENCE/DOCKET NUMBER: 24
          NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 92037-9103
                                                                                                                                                                                                                                                                                                                                          CITY: La Jolla
STATE: CA
                                                                                                                                                                                              APPLICATION NUMBER: US/09/287,682
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Matches:
Conservative:
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                                                                                                                                                                                                                                                                                                                                                                           Sequence 24, Application US/09287679 Patent No. 6258538 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                            APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
TITLE OF INVENTION: DNA Diagno
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLCGY: linear
MOLECULE TYPE: cDNA
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INFORMATION FOR SEQ ID NO: 24:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/6
FILING DATE: 18-MAR-1996
                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                         SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
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LENGTH: 252 base pairs
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CITY: I
                                                                                                                     COMPUTER: IBM CON
OPERATING SYSTEM:
                                          FILING DATE:
CLASSIFICATION:
                                                                           APPLICATION NUMBER:
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92037-9103
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: CA
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619-587-5360
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                                                                                                        Release #1.0, Version #1.25
              08/617,256
                                                                           US/09/287,679
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US-09-397-766-24
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: cDNA US-09-287-679-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alignment Scores:
                                                                                                                                                                                                                                                   Sequence 24, Application US/09397766 Patent No. 6268144
                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: K ster,
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (617)227-594 INFORMATION FOR SEQ ID NO:
SOFTWARE:
                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                          NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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                          COMPUTER: IBM CONTOPERATING SYSTEM:
                                                                                                    COUNTRY:
                                                                                                                                CITY: La Jolla
                                                                                                                                              STREET:
                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 GTGTGCGC-CGCCTGGTGCAGTACCGCGGCGAGGTGCAGGCCATGCTCGGCCAGAGCACC
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TELEFAX: 619-587-5360
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REGISTRATION NUMBER: 33,77
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ARE: PatentIn Rel APPLICATION DATA:
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(617)227-5941
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                                        IBM Compatible
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             #1.0, Version #1.25
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APPLICATION NUMBER:

us/09/397,766

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TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                         Sequence 24, Application US/09287681 Patent No. 6277573
                                                                                                                                          GENERAL INFORMATION: APPLICANT: K ster
                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White
STREET: 4250 Executive Square,
                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                           APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002I
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
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                             CITY: La Jolla
STATE: CA
COUNTRY: USA
ZIP: 92037-9103
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TELEFAX: 619-587-5360
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Query Match:
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TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 03 MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 2473
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEPHONE: 619-587-5360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION DATA:
FILING DATE: 18-MAR 1...
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
APPLICATION NUMBER: 03-MAR-1995
APPLICATION:
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OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Rel-
CURRENT APPLICATION DATA:
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                                                                                                                   No.:
   241
                 189 GluArgGlyLeu 192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 252 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Search completed: March 14, 2003, 20:18:27 Job time: 39.7343 secs

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MODEL-frame+_p2n.model -DEV=xlp
-Q-Ggn2_1/USPTO_Spool_VG09827854/runat_11032003_101612_27578/app_query.fasta_1.3576
-DB-Published_Applications_NA -QFMT-fastap -SUFFIX*p2n.rnpb -MINMATCH=0.1
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS-human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE-LOCAL -QUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=200000000 -USER-US99827854_@CGN_1 1_298_grunat_11032003_101612_27578
-NCPU=5 -ICPU=3 -NO_XLPXY -NO_MAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELEXT=7
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Match
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Ygapop 10.0 , X
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1585
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// Cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq: *
// Cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq: *
// Cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq: *
// Cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq: *
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US-09-870-759-129
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Sequence 129, App
Sequence 17, Appl
Sequence 8, Appli
Sequence 454, App
                                                                                                               Description
                                                                                                            ; FEATURE:
; NAME/KEY: CDS
; LOCATION: (61)...(1014)
; OTHER INFORMATION:
US-09-870-759-129
Percent Similarity:
                                               Pred.
                                                               Alignment Scores:
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US-09-870-759-129
                                                                                                                                                                                                                         SOFTWARE: Patent
SEQ ID NO 129
LENGTH: 1156
TYPE: DNA
ORGANISM: HOMO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: TERMAN, DAVID S
TITLE OF INVENTIO: COMPOSITIONS AND METHODS FOR TREATMENT OF NEOPLASTIC DISEASE
FILE REFERENCE: 870759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 129, Application US/09870759 Patent No. US20020177551A1
                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/208,128
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 166
                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/870,759
CURRENT FILING DATE: 2002-01-14
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Application
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GENERAL INFORMATION:
APPLICANT: Braun, Andreas
APPLICANT: Bonsal Aruna
APPLICANT: Kleyn Patrick
TITLE OF INVENTION: CARDIOVASCULAR DISEASE AND
CURRENT APPLICATION GENES AND POLYMORPHISMS ASS
TITLE OF INVENTION: CARDIOVASCULAR DISEASE AND
CURRENT APPLICATION UNMEER: US/09/802,640
CURRENT FILING DATE: 2001-03-09
NUMBER OF SEQ ID NOS: 122
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 1156
TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
NAME/KEY: CDS
LOCATION: (61)...(1014)
OTHER INFORMATION: Nucleotide Sequence encodin
OTHER INFORMATION: (APOE)
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Best Local Similarity:
Query Match:
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                                            AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
                                                      LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
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; ORGANISM: Homo
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Patent No. US20020123093A1
GENERAL INFORMATION:
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CURRENT FILLING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kypreos, Kyriakos E. TITLE OF INVENTION: Compounds and TITLE OF INVENTION: cholesterol FILE REFERENCE: 07180/004003
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NUMBER OF SEQ ID NOS: 850
SOFTWARE: PERL Program
SEQ ID NO 454
LENGTH: 1291
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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                                                                                                                               GENERAL INFORMATION:
                                                                                                                                      Sequence 454, Application Patent No. US20020137081A1
                                                                                                 APPLICANT: Olga Bandman
TITLE OF INVENTION: GENES D.
FILE REFERENCE: PA-0028 US
CURRENT APPLICATION NUMBER:
            NAME/KEY: misc_feato
OTHER INFORMATION:
NAME/KEY: unsure
LOCATION: 46
                                                                                            CURRENT FILING DATE:
      OTHER INFORMATION:
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods fo
TITLE OF INVENTION: Coholesterol levels wit
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
SOCTWARE: FastSEQ for Windows Version 4.0
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LENGTH: 1156
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; Patent No. US20020123099A1
; GENERAL INFORMATION:
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; ORGANISM: Homo
US-09-827-854-9
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TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Cohlesterol levels with FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cohlesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
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Patent No. US20020123093A1
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FILE REFERENCE: 689290-76

CURRENT APPLICATION NUMBER: US/09/954,456

CURRENT FILING DATE: 2001-09-18

PRIOR APPLICATION NUMBER: US/60/233,617

PRIOR APPLICATION NUMBER: US/60/234,052

PRIOR FILING DATE: 2000-09-20

PRIOR FILING DATE: 2000-09-20

PRIOR APPLICATION NUMBER: US/60/234,923

PRIOR FILING DATE: 2000-09-25

PRIOR APPLICATION NUMBER: US/60/235,134

PRIOR APPLICATION NUMBER: US/60/235,637

PRIOR APPLICATION NUMBER: US/60/235,637

PRIOR FILING DATE: 2000-09-26

PRIOR APPLICATION NUMBER: US/60/235,637

PRIOR APPLICATION NUMBER: US/60/235,720

PRIOR APPLICATION NUMBER: US/60/235,720

PRIOR FILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: US/60/235,720

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PRIOR APPLICATION NUMBER: US/60/235,863

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Patent No. US2
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TITLE OF INVENTION: Process
TITLE OF INVENTION: Sets
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APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
Alignment Scores Pred. No.: Score:
                                             ; FEATURE:
; OTHER INFORMATION:
US-09-880-107-2244
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US-09-880-107-2244
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APPLICANT: Kannis, Vassilis

APPLICANT: Kypreos, Kyriakos E.

APPLICANT: Kypreos, Kyriakos E.

TITLE OF INVENTION: Compounds and methods for lowering

TITLE OF INVENTION: Compounds and methods for lowering

TITLE OF INVENTION: Compounds and methods for lowering

TITLE OF INVENTION: Compounds and methods for lowering

TITLE OF INVENTION: Compounds and methods for lowering hypertriglyceridemia

FILLE OF INVENTION: Compounds and methods for lowering hypertriglyceridemia

CURRENT APPLICATION NUMBER: US/09/827,854

CURRENT FILING DATE: 2001-04-05

PRIOR APPLICATION NUMBER: US 09/679,088

PRIOR APPLICATION NUMBER: US 09/544,386

PRIOR FILING DATE: 2000-104-06

PRIOR FILING DATE: 2000-104-06
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SOFTWARE: FastSEQ for Windows Version
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; LENGTH: 1156
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                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing hypertriglyceridemia
FILE REFERENCE: 07180/004003
                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILLING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
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                                          SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLys
 CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAGGTCACCCAGGAACTGAGGGCG
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Alignment Pred. No. Score: Percent S	RESULT 12 US-09-967-013-5 Sequence 5, A FACENT NO. US GENERAL INFOR APPLICANT: APPLICANT: TITLE OF INV FILE REFEREN CURRENT APPLICURRENT APPLICURRENT APPLICURANT FILLING FRIOR APPLICOPATION FILLING FRIOR FILLING	Qy 301 Db 961	Qy 281 Db 901	Qy 26 Db 84	Qy 24 Db 78	Qy 22 Db 72	Qy 201 Db 661	Qy 181 Db 601	Qy 16 Db 54	Qy 14 Db 48	Qy 12 Db 42	Qy 101 Db 361	Оу 81 Оъ 301
t Scores: .: Similarity:	12  167-013-5  167-013-5  167-013-5  167-013-5  11 NO. US200200  ALINFORMATION  LOF INVENTION  REFERENCE: 11  RENT APPLICATION  RETLING DATE:  RETLING DATE:  REFLING DATE:			LeuGluGluGl 	1 SerArgThrAr              1 AGCCGGACCCG	1 LeuGlnGluAr               CTACAGGAGCG		ი – ი	1 LysLeuArgLy             1 AAGCTGCGTAA	1 GlnAlaMetLe               CAGGCCATGCT	1 ArgLeuGlyAl               1 CGGCTGGGCGC		
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Length: Matches: Conservative:	NETIC ANALYSIS OF REDEF	ValGlnAlaAlaValGlyThrSerAlaAlaProValProSerAspAsnHis 	SerTrpPheGluProLeuValGluAspMetGlnArgGlnTrpAlaGlyLeuValGluLy 	LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAlaArgLeuLys	SerArgThrArgAspArgLeuAspGluValLysGluGlnValA1aGluValArgAlaLys 	LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly 	ProLeuvalGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 	InAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaTleArgGluArgLeuGly 	LysLeuArgLysArgLeuLeuArgÅspAlaAspAspLeuGlnLysArgLeuAlaValTyr 	lnalametLeuGlyGlnSerThrĠluGluLeuArgValArgLeuAlaSerHisLeuArg            	rgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 	ThrProValAlaGluGluThrArgÅlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla 	LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu 
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			300	280 900	260 840	240 780	220 720	200	180	160 540	140	120	100 360

arity: 60.00% Indels: 194 10 Gaps: 194 AlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGl	CTCTGCTGGTTCTAGCTTCCTCTTCCCATTTCTGACTC	GAATTCTCTCTCAGCTTTGTCTCTCTCTCTCTCCTTCTGACTCAG	78	736 CGTCCTGGCTCTGTCTCTGTCCTTCCCTAGCTCTTTTATATAGAGACAGAGAG	78	796 CTCACTGTGTTGCCCAGGCTGGTCTTGAACTTCTGGGCTCAAGCGATCCTCCC	78	856 CCTCCCAAAGTGCTGGGATTAGAGGCATGAGCCACCTTGCCCGGCCTCCTAGCTC			916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTC	916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTC 78	916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTC 78	916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTC 78 976 GCCTCTGCCCCGTTCCTCTCTCCCTCTTGGGTCTCTCTC	916 TOGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTC  78 976 GCCTCTGCCCGTTCCTCTCTCTCTCTGGGTCTCTCTGGCTCA  78 036 CCCCATCCCAGCCCTTCTCCCCCGCCTCCCACTGTGCGACACCCTC	916 TOGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTC  78 976 GCCTCTGCCCCGTTCCTTCTCTCTCTCTTGGGTCTCTCTGGCTCA  78 036 CCCCATCCCAGCCCTTCTCTCCCCCCCTCTTGGGTCTCTCTC	16 TOGTCTCTGCCTCTGCCTCTGCATCTGCTCTCTGCATCTGTCTC  78  76 GCCTCTGCCCGTTCCTTCTCTCCCTCTTGGGTCTCTCTC	916 TCGTCTCTGCCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTCTCTT  78 976 GCCTCTGCCCGTTCCTTCTCTCTCCCTCTTGGGTCTCTCTGGCTCATCCCCATC  78 976 GCCTCTGCCCCGTTCCTTCTCCCCCCCCTCTTGGGTCTCTCTGGCTCATCCCCATC  78 97 GCCCATCCCAGCCCTTCTCCCCCGCCTCCCACTGTGCGACACCCTCCCCCCCTC  98 CCCCATCCCAGCCCTTCTCCCCCGCCTCCCACTGTGCGACACCCTCCCGCCCTC  99 CCCCATCCCAGCCCTTCTCCCCCGCCTCCCACTGTGCAAAGCACCTCCCGCCCTC  99 CCCCATCCCAGCCCTTCTCCCCCGCCTCCCACTGTGCAAAGCACCTCCCGCCCTC  99 CCCCATCCCAGCCCTTCTCCCCCGCCTCCCACTGTGCAAAGCACCTTCCGCAAATCGGAAACCGGAACCATGAAGGACCATGAAAGGAGTTGAAGGACCTACAAATCGGAAACCGGAAC  98 UGlnLeuThrProvalAlagluGluThrArgAlaArgLeuSerLysGluLeuG	916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTCCT  78  976 GCCTCTGCCCCGTTCCTTCTCTCCCTTTGGGTCTCTCTGGCTCATCCCCATC  78  79	916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCTCTCTCT	916 TCGTCTCTGCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTCTTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTC	916 TCGTCTCTGCCCTCTGCCCTCTGCATCTGCTCTCTGCATCTGTCTCTCTC
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APPLICANT: Rosen et al.

TITLE OF INVENTION: Nucleic Acids, Proteins;

FILE REFERENCE: PAL04

CURRENT APPLICATION NUMBER: US/09/925,302

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: PCT/US00/05918

PRIOR FILING DATE: 2000-03-08

PRIOR FILING DATE: 1909-03-12

NUMBER OF SEQ ID NOS: 896

SOFTWARE: PATENTIN VET. 2.0

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                                                                                                                                                                                                                           Alignment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 133, Application US/09925302 Patent No. US20020044941A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
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                                                     CGCCTGGTGCAGTACCGCGGCGAGGTGCASGCCATGCTCGGCCAGAGCACCGAGGAGCTG
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                                                            Percent Similarity:
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Query Match:
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; OTHER INFORMATION: n=
US-09-964-824A-374
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                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. US20020102531A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                             SOFTWARE: PatentIn version SEQ ID NO 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 374,
                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Cancer TITLE OF INVENTION: Sets FILE REFERENCE: 689290-73
                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/964,824A CURRENT FILING DATE: 2001-09-27
                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US/60/236,033 PRIOR FILING DATE: 2000-09-28
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PRIOR FILING DATE: 2000-09-28
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                                                                                                                                                                                PEATURE: misc_feature
                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                      TYPE: DNA
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APPLICANT: Vockley, Joseph G.

APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.

TITLE OF INVENTION: Gene Expression Profiles
FILE REFERENCE: 44921-5028-W0
CURRENT APPLICATION NUMBER: US(09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
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                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
3EQ ID NO 2491
LENGTH: 478
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OTHER INFORMATION: n = a or c or g or
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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Ygapop 10.0 , Ygapext
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## SUMMARIES

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## ALIGNMENTS

RESULT 1
B1670367
LOCUS
DEFINITION
MRNA Sequence.

ACCESSION
B1670367.1 GI:15584600
KEYMORDS
SOURCE Homo sapiens
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
AUTHORS
NATIONAL TO BEST 12-SEP-2001
REPERENCE 1 (bases 1 to 1027)
AUTHORS
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GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI)
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNI
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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/db_xref="taxon:9606"
/clone="IMAGE:5312024"
/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA sequence.
BG763371
BG763371.1 GI:
EST.
                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTPDTP
Tissue Procurement: ATCC/DCTPDTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution: MGC clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.lini.gov
plate: LLCM1718 row: b column: 10
High quality sequence stop: 817.
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (Dases 1 to 842)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
                                                                                                                                                                                                                                                                                                               Unpublished (1999)
Contact: Robert Strausberg,
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/tissue_type="melanotic melanoma,
/lab_host="DH10B (phage-resistant
                                                      /clone="IMAGE:4860585"
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/db_xref="taxon:9606"
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              ySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
                                                                     LeuGlnGluArgAlaGlnAlaTrpGlýGluArgLeuArgAlaArgMetGluGluMet-Gl
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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BI668318
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Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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/Clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium ()
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution informatic
                                                          1 (bases 1 to 938)
NHH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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/lab_host="DHIOB (phage-resistant)"
/note="Organ: skin; Vector: PDFI; Site_1: XhoI; Site_2:
RCORI; cDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GCCACGAAC(G). Size-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
II RT (Life Technologies). Note: this is a NIH_MGC
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NIH-MGC http://mgc.nci.nih.gov/
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603249241F1 NIH_MGC_96
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                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone='TMAGE:5301010"
/clone='Ib="NIH_MGC_96"
/tissue_type="Nypothalamus"
/lab_host="DH10B"
                                                                            D
                                                                        /note="Organ: brain; Vector: pBluescript R (modified pBluescript KS+|); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIHH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library." a 284 c 361 g 120 t
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ACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
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1 (bases 1 to 800)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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Plate: LLCM1875 row: i column: 03
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/lab_host="PH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; cDNA made by oligo-dT priming. Directionally cloned
into EcORI/XhoI sites using the following 5' adaptor:
GGCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
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/db_xref="taxon:9606"
/clone="IMAGE:5420618"
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM1419 row: k column: 08
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Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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/lab_nost="Norgan: eye; Site_1: XhoI; Site_
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                                                                                                        etGlySerArg-ThrArgAspArgLeuAspGluValLysGluGlnValAlaGlu-ValAr 258
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                                                                                                                                                                                                                                                                                       LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                                                                                                            --GlyProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyG
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                                                                                                                                                                                                                    CAGGGCCGGGGCCCGAGGGCCGCGGCCCTCAAGCGCCATCCGCGAGCGCCTT
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                                                                                                                                                                       GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                               CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCAGGTCACCCAGGAACTGAGGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 922)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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BI597743
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcg: ); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carnincl, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library." a 273 c 359 g 114 t
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/db_xref="taxon:9606"
/clone="IMAGE:5300259"
/clone_lib="NIH_MGC_96"
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Mismatches:
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                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1612 row: e column: 20
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National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                             mRNA sequence.
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                                                                                 quality sequence stop:
Location/Qualifiers
/clone_lib="NIH_MGC_40"
/tissue_type="carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4760827"
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/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."

148 a 242 c 324 g 103 t
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Alignmer Pred. No Score: Percent Best Loc Query Ma	ment Scores:       3.5e-115       Length:       817         No.:       1190.50       Matches:       252         ent Similarity:       97.32%       Conservative:       2         Local Similarity:       96.55%       Mismatches:       5         Local Similarity:       75.11%       Indels:       4         75.11%       Gaps:       0
us-09-	-827-854-15 (1-317) x BG774871 (1-817)
9 57	1 MetLysValleuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 20 
9 54	21 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 40
₽ 2 <sub>4</sub>	41 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 60 
ρ Σ	61 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80
8 8	81 LeuMetaspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluClnLeu 100 
Dy Ob	101 ThrprovalAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla 120
9 54	121 ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140
₽ ¥	141 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160
9 Y	161 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr 180
8 8	181 GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly 200
β <del>2</del> 4	201 ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro 220
Dy Ob	221 LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetG 240
ου Σ	240 lySerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAla 259           ::

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CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
                                                       GGCCAGCGCTGGGAACTGGGACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                                   GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.lln1.gov
plate: LLAM11726 row: g column: 10
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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National Institutes of Health, Mammalian
Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:5287329"
/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov Plate: LLCM1790 row: g column: 17 High quality sequence stop: 829. Location/Qualifiers
                                                                                                                                                                                                                                                   Contact: Robert Strausberg, | Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                             NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          602763768F1 NIH_MGC_42 Homo sapiens cDNA clone IMAGE:4899112
                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                         cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.B. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                    human
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/db_xref="taxon:9606"
/clone="IMAGE:4899112"
/clone_lib="NIH_MGC_42"
/tissue_type="epithelioid carcinoma cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1:
                                                                                                       1. .845
                                                                                        /organism="Homo sapiens"
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                                                                                                     ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
                                                                                                                                   SerArgThr-ArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLy
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  AGCCGGACCGCGA--
                                             CTACAGGAGCGGGCCAGGCTG---
                                                                                         CCCCTGGTGGAACAGGGCCGCGTGCGGGCCGC-ACTGTGGGCTCCCTTGGCGGCAG-CCG
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Directionally cloned into ECORI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Ling
Hong in the laboratory of Gerald M. Rubin (University of
California, BerkeLey) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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                                             -GGGCGAGCGGTGCGCGCGCGGATGGAGGAGATGGGC
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TITLE
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193
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                                                                                   GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
                                                                 GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGCAGACCGAGTGGCAGAGC
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BG707147
BG707147.1 GI:13983201
EST.
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LiAM10672 row: e column: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.I
cDNA Library Preparation: Michael J. Bro
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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602670283F1 NIH_MGC_96 Homo
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="hypothalamus"
/lab_host="D1018"
/lab_host="D1018"
/note="Organ: brain; Vector: pBluescriptR (modified /note="Organ: brain; Vector: pBluescriptR (sal1.xhoI (gtcgag pBluescript KS+); Site_1: BamHI; Site_2: Sal1.xhoI (gtcgag pBluescript KS+); Site_1: BamHI; Site_2: Sal1.xhoI (gtcgag ); Oligo-dr primed using primer 5'-TTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninc1, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."
43 a 234 c 311 g 102 t
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/clone="IMAGE:4793084"
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                                                                         DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM11691 row: 1 column: 04 high quality sequence stop: 812.
                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consorti
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Mammalia; Eutheria; Primates; Ca
1 (bases 1 to 919)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, N
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Contact: Robert Strausberg, Ph.D
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/organism="Homo sapiens"
/db_xref="taxon;9606"
/clone="IMAGE:5274003"
/clone_lib="NHH_MGC_95"
/tissue_type="hippocampus"
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/note="Organ: brain; Vector: pBluescript (modified pBluescript KS+); Site\_1: BamHI; Site\_2: SalI-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTVN-3', size-selected for average insert size 2.5 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH\_MGC Library."

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Best Local Similarity:
Query Match:
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CAGCCGGACCCGGGACG - - CTGGACGAAGTGAAGAGCACGTGGGCCGAAGGG - TGCGCCAA
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GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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Tissue Procurement: DCTD/DTP
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                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incytte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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602669093F1 NIH_MGC_96 Homo sapiens
                                                                                                                                                                      found through the I.M.A.G.E. Consortium/LLNL
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene
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/clone_lib="NIH_MGC_96"
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Command line parameters:

MODEL-frame+_p2n.model -DEV-xlp
-Q-/cgn2_1/USPTO_spool/US09827854/runat_11032003_101610_27486/app_query.fasta_1.3576
-DB-GenEmbl -QFMT-fastap -SUFFIX-p2n.rge -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXILEN-200000000
-USER-US09827854_@CGN_1_1_13008_@runat_11032003_101610_27486 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MAAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120
-WARN_TIMEOUT-30 -THREADS-1 -XGAPOP+10 -XGAPEXT-7
-YGAPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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PI IKEDA YASU
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strandedness: Double;
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hypothetical: No;
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Key Location/Qualifiers
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RS Teranishi,Y., Matsui,Y., Ikeda,Y. and Kimura,M.

RRS Teranishi,Y., Matsui,Y. and Kimura,M.

PRODUCTION OF HUMAN APOLIPOPROTEIN E-LIKE PROTEIN

ALL PROTEIN IND LTD

OS Human (Homo sapiens)

PN JP 1986096997-A/1

PD 15-MAY-1986

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

PF 16-OCT-1984 JP 1984216987

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/db_xref="taxon:32644"
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Sequence 63
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                                                        Morin, P.J., Sherman-Baust, C.A., Pize Tumor markers in ovarian cancer Patent: WO 017517-A 63 jl-0cT-2001; THE SECRETARY OF THE DEPARTMENT OF H Location/Qualifiers
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Eukaryota; M
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                     /organism="Homo sapiens"
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BD004278
Fujita, S., Hamanaka, H., Fukui, Y. and Y
Apo E humanized mammal
Patent: JP 2001017028-A 2 23-JAN-2001;
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/2
PD 23-JAN-2001
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                                                               Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
Mammalia; Eutheria; Primates;
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JP 2001017028-A/2.
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(6) sites; mutations resulting in type III hyperlipoproteinemia. Apo E is a component of normal human very low density lipoprotein. There are six human apo E phenotypes known to result from a single structural gene, three of the common alleles being epsilon-4, epsilon-3 and epsilon-2. This sequence appears to be of the epsilon-3 allele. [1] argues that the apo E polymorphism involves mutations in the structural coding region; for example the epsilon-2 phenotype which is characterized by hyperlipoproteinemia is thought to result from a c to t change (arg to cys) at base 586 below [3], [5]. The sequence shown is 57% homologous with human apo A-I and 81% homologous with rat apo E. For the epsilon-4 sequence,
                                                                                                                                                                                                                                                                                                                    J. Clin.
89198059
                                                                                                                                                                                                                                                                                                                                                                        Rall, S.C. Jr., Newhouse, Y.M., Clarke, H.R., Weisgraber, K.H., McCarthy, B.J., Mahley, R.W. and Bersot, T.P.
Type III hyperlipoproteinemia associated with apolipoprotein phenotype E3/3. Structure and genetics of an apolipoprotein E
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Gill.L.L., Peoples, O.P., Pearston, D.H., Robertson, F.W., Humphries, S.E., Cumming, A.M. and Hardman, N. Isolation and characterisation of a variant allele of thuman apolipoprotein E Biochem. Biophys. Res. Commun. 130 (3), 1261-1266 (1985)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 (bases 250 to 777), Gill,L., M Wallis,S.C., Rogne,S., Gill,L., M Williamson,R. and Humphries,S. The isolation of cDNA clones for detection of apoE RNA in hepatic EMBO J. 2 (12), 2369-2373 (1983)
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83082756
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                                                                                                                                                                                                                                                                                                                                                      variant
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11422-11422 (1983)] correction of [1]. [4
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and [3].
Apo E is located on chromosome 19 --Jackson, Bruns and Breslow,
PNAS USA 81, 2945-2949 (1984) -- and is believed to be linked to
the Apo C-II gene (see separate entry).
The two mutations causing type III hyperlipoproteinemia
(apolipoprotein E phenotype E3/3) produces substitutions of Arg for
Cys at amino acid 112 and Cys for Arg at amino acid position 142.
Draft entry and printed copy of sequence for [1] kindly provided by
L.L. Gill, University of Aberdeen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          See the Separate [J. Biol. Chem. 258, 11422-11422 (1700), --
relative to [1] to record the
revisions in the FEATURES table. The sequence below is
Biol. Chem. 258, 11422-11422 (1983)]
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                                                                                                                      CGGCTGGGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
                 ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140
                                                                                                                                                                                     GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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                                                                           ThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla
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/replace="g"
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                                                                                                                                                                                                                                                                                                        Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvanah Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jili Vardy, George Yang, Scott Zuyderduyn, Marco Marra.
                                                                                                                                                                                                               Series: IRAL Plate: 6 Row: h Column: 23
This clone was selected for full length sequencing becaused the following selection criteria: matched mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (20-FEB-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          complete cds.
BC003557
                                                                                                                                                                                                                                                               Clone distribution: MGC clone distribution through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Genome Sequence Centre,
DLA Sequency, Vancouver, BC, Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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                                              /note="Vector:
46. .999
                                                                          /tissue_type="Eye, retinoblastoma"
/clone_lib="NIH_MGC_16"
/lab_host="DH10B-R"
/codon_start=1
/product="apolipoprotein
/protein_id="AAH03557.1"
                                                                                                                       /db_xref="taxon:9606"
/clone="MGC:1571 IMAGE:3355712"
                                                                                                                                                       /db_xref="LocusID:348"
                                                                                                                                                                       /organism="Homo sapiens"
                                                                                                                                                                                                     Location/Qualifiers
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Percent Similarity:
Best Local Similarity:
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E08423 1110 bp
B08423 phuman apolipoprotein
E08423 GI:2176540
JP 1994315392-A/1.
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ELALGRFWYLKRWVQTLSEQYQEELLSSQVTQELRALMDETMKELKAYKSELEEBQLTP
VAEETRARLSKELQAAQARLGADMEDVCGRLVQYRGEVQAMLGQSTEELRVRLASHLR
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QPLQERAQAWGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ
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GlnAlaMetLeuGlyGlnSerThrGlüGluLeuArgValArgLeuAlaSerHisLeuArg
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METHOD FOR PRODUCING ADOLLPOROTEIN
PATENT: JP 1994315392-A 1 15-NOV-1994;
MITSUBISHI KASEI CORP
OS HOMO Sapiens (human)
PN JP 1994315392-A/1
PD 15-NOV-1994
PF 11-JUN-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF 11-JUN-1985 JP 1994015433
PF (12P21/02,CO7K13/00,C12N5/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/10,C12N15/
                                                                                           ArgLeuGlyAlaAspMetGluAspVaiCysGlyArgLeuValGlnTyrArgGlyGluVal 140
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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MORIMOTO HIRONORI, TERANISHI YUTAKA
C12P21/02,C07K13/00,C12N5/10,C12N15/12,C12N15/18,(C12P21/02,
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/db_xref="taxon:9606"
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3 Fujita, S., Hamanaka, H., Fukui, Y. and Yokoyama, M.
Apo E humanized mammal
- Patent: JP 2001017028-A 1 23-JAN-2001;
MITSUBISHI CHEMICAL CORP
OS Homo sapiens (human)
PN JP 2001017028-A/1
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128919
PR
GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                                                                                                       ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
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/db_xref="taxon:9606"
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Location/Qualifiers
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Attie, A.D., Gretch, D.G., Sturley, S.L.
Production of recombinant proteins in
Patent: US 5472858 A 1 05-DEC-1995;
Location/Qualifiers
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J. Biol. Chem.
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M12529.1 GI:178848
apolipoprotein.
Homo sapiens (clone:
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McLean,J.W., Elshourbagy,N.A., Chang,D.J., Mahley,R.
                                                                                                                                                                   Human apolipoprotein
                                                                                                                                                                              Taylor, J.M.
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                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/map="19q13.2"
         /gene="APOE"
<1. .1157
                                                                                                          Location/Qualifiers
1. .1157
/gene="APOE"
                             /dev_stage="57-year
1. .1157
                                              /tissue_type="liver"
                                                         /clone="pHAE[112,178,813]."
/sex="male"
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/replace="c"
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/replace="g"
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/product="apolipoprotein
/note="G00-119-691"
370
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/protein_Id="AAB59518.1"
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/db_xref="GDB:G00-119-691"
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VAEETRARLSKELQTAQARLGADMEDVCGRLVQYRGEVQAMLGQSTEELRVRLASHLR
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/note="g in pHAE[112,178];
/replace="a"
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/note="a in pHAE[112,178]; g
/replace="g"
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/note="precursor"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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JP 2001017028-A/3.
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Sequence 2
A62340
A62340.1
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Van, L.F., Burbach, J.P. and Grosveld, F.
DIACNOSIS METHOD AND REAGENTS
Patent: WO 9712992-A 30 10-APR-1997;
POYAL NETHERLANDS ACADEMY OF A (NL)
Other publication AU 7142796 970428.
Location/Qualifiers
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-LOOPEXT=0_-UNITS=bits_-START=1 -END=-1 -MATRIX=biosum62_-TRANS=human40_cdi
-LIST=45_-DOCALIGN=200_-THR_SCORE=pct_-THR_MAX=100_-THR_MIN=0_-ALIGN=15_-CMINMODE=LOCAL_-OUTEMT=pto_-NORM=ext_-HEAPSIZE=-900_-MINLEN=0_-000000000
-USER=US09827854_eCGN_1_1_1201_erunat_1103200_101609_27476_-NCPU=6_-TGU=3
-NO_XLPXY_-NO_MMAP_-LARGEQUERY_-NEG_SCORES=0_-WAIT_-LONGLOG_-DEV_TIMEOUT=120
-WARN_TIMEOUT=30_-THREADS=1_-XGAPOP=10_-XGAPEXT=0.5_-FGAPOP=6_-FGAPEXT=7
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2: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT:*

3: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1982.DAT:*

4: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1983.DAT:*

5: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT:*

6: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT:*

7: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT:*

8: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1986.DAT:*

9: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1989.DAT:*

10: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1999.DAT:*

11: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1991.DAT:*

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13: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1991.DAT:*

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Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being part and is derived by analysis of the total score distribution. printed D

## SUMMARIES

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	Human apolipoprote	~	Human secreted pro	ApoE4L protease cD	Human apolipoprote	ApoE4 cD	Human ApoE4Lx2 cDN	ApoE4Lx2 protease	Human diagnostic a	Thyroid cancer rel	adenocarc		Himan CDNA for now	Nucleotide sequenc	Human cDNA encodin		alpha.	OD.	apolip	Apo E		apol		Human apolipoprote	w .			Human ApoE4 coding	Sequence encoding	- 0		- 5		Human apolipoprote		Human apolipoprote			Human ApoE3 coding	Apolipoprotein E o	Human apolipoprote	Human apolipoprote	scription

## ALIGNMENTS

RESULT 1 AAD26035 Human apolipoprotein E (APOE) cDNA. 26-MAR-2002 (first entry) AAD26035; AAD26035 standard; cDNA; 954

Human; antilipaemic; neuroprotective; nootropic; genetic variant; APOE; apolipoprotein E; haplotyping; familial dysbetalipoproteinaemia; therapy; genotyping; type III hyperlipoproteinaemia; Alzheimer's disease; SNP; atherosclerosis; single nucleotide polymorphism; chromosome 19q13.2; ss. Homo sapiens.

Location/Qualifiers

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Alignment
Pred. No.:
                       Percent Similarity:
Best Local Similarity:
                                                                                                 Query Match:
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                                                                                                                               The patent discloses novel genetic variants of human apolipoprotein CE (APOE) gene. The invention also relates to compositions and methods of for haplotyping and/or genotyping the APOE gene. The haplotyping methods of the invention are useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with APOE activity, e.g. can it is a dispersion of the invention are useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with APOE activity, e.g. can it is a candidate agent for treating a specific condition or disease predicted to be associated with APOE activity and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with APOE activity and in the design of cor disease predicted to be associated with APOE activity. Genotyping APOE to treat a specific condition or disease associated with APOE activity. The present sequence is a cDNA encoding human APOE protein.
                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                            Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene -
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                                                                                                                 LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                                                                                                                  GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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                                                                                                                                                                                                                                                                         ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
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DB; AAP60507.
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CC The invention relates to methods for diagnosing and prognosing ovarian CC tumours in an individual via the detection and measurement of the CC expression of ovarian tumour marker genes (ABA83081-ABA83122, ABA83180, CC ABA83182 and ABA83184) or segments thereof (ABA83181-ABA83129, ABA83179, CC ABA83182 and ABA83183). The methods of the invention are useful for CC detecting an ovarian tumour in a patient, for identifying an individual CC at increased risk for developing ovarian cancer, in prognostic tests for CC monitoring disease status in a patient being treated for ovarian cancer. The methods can additionally be used to identify a particular tumour as CC being an ovarian tumour (i.e., an epithelial ovarian tumour selected from Serous cystadenoma, borderline serous tumour, serous cystadenocarcinoma, come cipstadenocarcinoma, andometrioid carcinoma, undifferentiated carcinoma, CC cystadenocarcinoma, pendometrioid carcinoma, undifferentiated carcinoma, CC cystadenocarcinoma, cystadenofibroma, adenofibroma and Brenner CC using SAGE (serial analysis of gene expression) and were found to be coverexpressed in a broad variety of ovarian epithelial tumour cells call implicated in immune response pathways, in the regulation of cell proliferation and in protein folding, and many of these are membrane-CC implicated in ambure response pathways, in the regulation of cell prognostic markers, the ovarian tumour marker genes or their encoded CC proteins may be used as therapeutic targets for the treatment and CC prevention of ovarian cancer. Sequences ABA83081-ABA83182, ABA83180, ABA83184 represent the ovarian tumour marker genes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mucinous cystadenoma; borderline mucinous tumour; endometriold carcinoma; undifferentiated carcinoma; clear cell adenocarcinoma; cystadenofibroma; adenofibroma; Brenner tumour; serial analysis of gene expression; SAGE; immune response pathway; cell proliferation regulation; protein folding; membrane localised; secreted; therapeutic target; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting and identifying ovarian tumor, identifying increased risk for developing ovarian cancer, and determining effectiveness of ovarian cancer treatment, by measuring expression level of ovarian tumor marker
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P-PSDB; ABB50287.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epithelial tumour; cancer; diagnosis; prognosis; disease moni identification; serous cystadenoma; borderline serous tumour;
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                                                                                                                                              The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammaal lacking an endogenous,
                                                                                                                                                                                                                                                                                                                                                                                               (KOSP-)
                                                                                                     hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein receptor or is at risk for developing atherosclerosis due accumulation of lipoprotein remnants in the bloodstream or
                                                                                                                                                                                                                                      Claim 14;
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immunomodulatory; cytostatic; neuroprotective; vulnerary; nootropic; anticonvulsant; antiarthritic; cerebroprotective; antifungal; antiviral antibecterial; antiallergic; dermatological; haemostatic; antiasthmatic thrombolytic; immunogen; antibody; gene therapy; neurological disorder; parkinson's disease; inflammatory disorder; cancer; asthma; osteoporosi
                                                                                                                                             Human cDNA encoding a novel human
                                                                                                      Human; novel protein; ss; Antianaemic; osteopathic;
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(UYBO-)
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US-09-827-854-15_COPY_1_203 (1-203) x AAF84314 (1-1156)
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Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to an ApoE hun present sequence is the coding sequence for in the method of the present invention. The can be used for screening for agents useful Alzheimer's disease and arteriosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for treating arteriosclerosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1156 BP; 208 A; 367 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 11-12; 22pp; Japanese.
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                                        LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
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05-APR-2001;
                                                                      New apolipoprotein E polypeptide and nucleic acid, useful for lowering cholesterol, delaying the onset of or treating atherosclerosis in
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US-09-827-854-15_COPY_1_203 (1-203)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             isoprotein, apoE2 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertridglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                                                                  ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                                                                             LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                                                                                                                                                                                                                                                          CGGCTGGGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
                                                                                                                                                                                   ACCCCGGTGGCGGAGGAGACGCGGCACGGCTGTCCAAGGAGCTGCAGGCGCGCAGGCC
                                                                                                                                                                                                                               CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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ВP

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Alignment
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                                                                                                              Best Local Similarity: Query Match:
DB:
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                                                                                            US-09-827-854-15_COPY_1_203 (1-203) x AAD22051 (1-1156)
                                                                                                                                       Percent Similarity:
                                                                                                                                                Score:
                                                                                                                                                                                                The present sequence is a human apolipoprotein E (apoE) allele, apoE2* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LI receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or had defect in remnant removal.
                                                                                                                                                                                                                                                                                    Claim 14;
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04-OCT-2000; 2000US-0679088
05-APR-2001; 2001US-0827854
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                                                                                                                                                                                                                                                                                                 apolipoprotein E polypeptide and nucleic acid, useful for lowering lesterol, delaying the onset of or treating atherosclerosis in mal, without inducing hypertriglyceridemia -
GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                                    MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal
                                   GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
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                         CAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGCAGACCGAGTGGCAGAGC
                                                           ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
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pw density lipoprotein; LDL; ds.
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WPI; 1996-029812/03
P-PSDB; AAR86791.
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                                                                                                                                                                                                                                                                                                                               Misc_feature
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                Recombinant; human; apolipoprotein-E; ApoE; insect cells; larva; Manduca sexta; Autographica californica nuclear polyhedrosis virus; haemolymph; lipid complex; biologically active; ds.
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nuclear
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                                                                       CAGGCCGGGGCCCGAGGGCGCCGAGCGCCTCAGCGCCATCCGCGAGCGCCTGGGG
                                                                                                                      AAGCTGCGTAAGCGGCTCCTCCGCGATCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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                                                                                                                                                                                                                                                                                                                               progression of liver cancer, hepatcoellular carcinoma or metastatic liver tumour in a patient, and differentiating metastatic liver cancer from hepatocellular carcinoma in a patient, involving detecting the level of expression of two or more genes represented in ABN93503-ABN97455 in a tissue sample. The method of the invention has hepatotropic, and cytostatic activity. The method is useful for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma and metastatic liver carcinoma in a patient. The method is useful for identifying expression profiles which serve as useful diagnostic markers as well as markers that can be used to monitor disease states, disease progression, drug toxicity, drug efficacy and drug metabolism.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                           Sequence 1157
                                                                                                                                                                                                                                                                                                                     at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1;
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                                Disclosure;
                                                                      Identifying drugs for and diagnosing benign prostatic hyperplasia, by detecting expression levels of one or more genes in prostate cells from patient that are differentially regulated compared to normal prostate
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05-JUN-2001;
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cc detecting the expression levels of one or more genes in prostate cells cfrom the subject that are differentially regulated compared to normal compared tells (II) comprises preparing a first gene expression profile cof BPH cells or BPH-like cell population, exposing the cells to the agent preparing a second gene expression profile of the agent exposed cc cells, and comparing the first and second gene expression profiles. (Cl) is useful for diagnosing the onset or progression of BPH. (II) is cusful for identifying an agent that modulates the onset or progression cc of BPH. The methods are useful to present information identifying the expression cross of the expression level in a tissue or cells, by comparing the expression cc level of genes given in the specification in the tissue or cells to the capters of the expression of gene in the database, and displaying the cc expression the elevel of the expression level in BPH. Agents using (II) are useful for treating BPH or prostate cancer. ABK641860 represent human cross that the time tells ample content of the prostate cancer. ABK641860 represent human cross that the treating BPH or prostate cancer. ABK641860 represent human cross that the treating BPH or prostate cancer. ABK641860 represent human cross that the treating BPH or prostate cancer.
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prostatic
hyperplasia gene sequences of the invention.
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Sequence 1157 BP; 212 A; 370 ç; 426 G; 149 Τ; 0 other;

1.07e-86 1005.00 99.01% 99.01% 99.11%

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ng; prostate; pancreas; carcinoma; antitumour; cancerous;
gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;
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Alignment Scores:
                                                                                                                      The present invention describes a method (MI) for screening for an CC anti-neoplastic agent. The method involves exposing cells to a chemical CC agent to be tested for anti-neoplastic activity, determining a change in CC expression of at least one gene (I) of a signature gene set, where (I) CC comprises a sequence (S) selected from 8447 sequences (given in ABL6164 CC to ABL70110), or is at least 95% identical to (S), where a change in CC expression is indicative of anti-neoplastic activity. (I) has cytostatic CC activity and can be used in gene therapy. MI can be used for screening CC an anti-neoplastic agent, and can be used for product which CC is the data collected with respect to the anti-neoplastic agent as a CC result of MI, and the data is sufficient to convey the chemical CC treatment of cancer such as colon, breast, stomach, lung, thyroid, CC canonicar contain, kidney, prostate or pancreatic cancer.
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Soppet DR,
                                                                      adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer, infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.
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                                   Sequence 1157
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Matches:
Conservative:
Mismatches:
Indels:
Gaps: 1157 201 0 2 0 0 541 140 421 120 361 301 241 60 40 121 160 481 100 181 20

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DB; AAP51204.
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Search completed: March 14, 2003, 12:43:05 
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-MODEL=frame+_p2n.model -DEV=x1p
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-DB=Issued_Patents_Na -OpMT=fastap| -SUFFIX=p2n.rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=+Dits -START=1 -END=-1 -MATRIX=Dlosum62 -TRANS=human40.cd1
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
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-USER=USO9827854_@CGN_1_1_224_@rumat_11032003_101611_27510 -NCPU=6 -ICPU=3
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-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq: *
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3: /cgn2_6/ptodata/2/ina/6A_COMB.seq: *
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STATE: WI
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ZIP: 53701-2113
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## ALIGNMENTS

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US-07-709-949-1
Sequence 1, Application US/07709949
Sequence 1, Application US/07709949
Sequence 1, Application US/07709949
Sequence 1, Application US/07709949
Sequence 1, Application (No. 5472858)
GENERAL INFORMATION:
APPLICANT: Attie, Alan D
APPLICANT: Sturley, Stephen L
APPLICANT: Sturley, Stephen L
APPLICANT: Sturley, Stephen L
APPLICANT: Sturley, Stephen L
APPLICANT: Beckage, Nancy E
TITLE OF INVENTION: Insect Larvae
NUMBER OF SEQUENCES: 2
CORRESSPONDENCE ADDRESS:
ADDRESSEE: Quarles & Brady
STREET: P.O. Box 2113
COUNTRY: USA
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/07/709,949
FILING DATE: 19910604
CLASSIFICATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 9629691801
TELEPHONE: (608)251-5000
TELEPAX: (608)251-5000
TELEPAX: (608)251-9166
INFORMATION FOR SEQ ID NO: 1:
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SEQUENCE CHARACTERISTICS: LENGTH: 1157 base pairs

TYPE: NUCLEIC ACID STRANDEDNESS: doub

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Best Local Similarity:
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ORIGINAL SOURCE:
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PUBLICATION INFORMATION:
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JOURNAL: Journal of Biological Chemistry
VOLUME: 259
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DATE: 1984
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                          AGCTGCGTAAGCGGCTCCTCCGCGATCCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/
FILING DATE: 02-Oct-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 95/
FILING DATE: 02-Oct-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
FILING DATE: 01-Jan-1996
ATTORNBY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 660 base pairs
TYPE: nucleic acid
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NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: van Leeuwen, Frederik Willem
APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
                                  122 GAGCAAGCGGTGGAGACAGAGCCGGAGCCCGAGCTGCGCCAGCAGACCGAGTGGCAGAGC
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41 GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
                                                 21 GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer
                                                                                                62 ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 96,048-A (3255/00784)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette, 3.50 inch, COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDPERFECT 6.1
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                                                                                                                                                                CURRENT APPLICATION NUMBER: US/08/949,155
APPLICATION NUMBER: US/08/949,155
FILING DATE: CONCURRENTLY Herewith
CLASSIFICATION: 800
PRIOR APPLICATION NUMBER: US 60/027,338
FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:

APPLICATION DATA:

11-OCT-1996
PRIOR APPLICATION DATA:
                          TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Compositions and Methods TITLE OF INVENTION: Generation of Transgenic NUMBER OF SEQUENCES: 51
                SEQUENCE CHARACTERISTICS:
                                                                    REFERENCE/DOCKET NUMBER: TI
                                                                                                                          APPLICATION NUMBER: US 60 FILING DATE: 09-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Piedrahita, Jorge APPLICANT: Bazer, Fuller W
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                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                               NAME: Hibler, David W. REGISTRATION NUMBER: 4
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                                                                                Sequence 5, Application Patent No. 6369294
GENERAL INFORMATION:
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LOCATION:
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             NUMBER OF SEQUENCES:
                         TITLE OF INVENTION: Compositions and Methods
Generation of Transgenic
                                                                 APPLICANT: Piedrahita,
                                                                                                                                                                    AlaValTyrGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGlu
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    No.:
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                                                                                                                                                                                                159 TGGCAGGGCAGCCCTGGGAGCAGGCCCTGGGCCGCTTCTGGGATTACCTGCGCTGG
 339 GCGCAGCTGGGCCCCGTGACCCCAGGAGACGCAGGCGCCCTGTCCAAGGAGCTGCAGGCG
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                                                                                78 LeuArgAlaLeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGlu
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               98 GluGlnLeuThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAla
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LOCATION: 51..1
SEQUENCE DESCRIPTION:
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FILING DATE: <UNKNOWN>
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
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MEDIUM TYPE: Floppy disk
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LOCATION:
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TYPE: nucleic acid
STRANDEDNESS: single
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FILING DATE: 28-MAI-2001
CLASSIFICATION: CURKNOWN>
APPLICATION DATA:
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TELEFAX: (713) 789-2679
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Gaps:
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Best Local Similarity:
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Patent No. 6271436
                                                                                                                                            TELEFAX: (713) 789-2679
INFORMATION FOR SEO ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                      TELEPHONE: (512) 418-3000
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CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE AND DURKEE
STREET: P.O. Box 4433
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                                                            No.:
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CITY: H
                                                                                                                  TOPOLOGY:
                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK:177
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/027,338 FILING DATE: 11-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: Concur
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ZIP: 77210-4433
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                             AGCGCCTCCGCGAGCGCCTCGGGCCCCTGGTG 3409
                                            SerAlaIleArgGluArgLeuGlyProLeuVal 203
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                                                                                                               LeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeu 192
                                                                                                                                             AGCCGCCTGGCTTCCCACCTGCGCAAGCTGCGCAAGCGGCTGCTCCGCGACACCGAGGAC
                                                                                                                                                                                                    TTGGTGCTCTACCGCAGCGAGGTGCACAACATGTTGGGCCCAGACCACCGAGGAGCTGCGG
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                                                                                                                                                                                                                                                                                                                                                                                                       ----ArgAlaLeuMetAspGluThrMetLysGluLeuLysAlaTyr 92
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                                                                                                                                                                                                                                                                                                                Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                               Score:
                                                                                                                                                                                                                                                                                                                                                           Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                     ; SEQUENCE DESCRIPTION: SEQ US-09-819-964-51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 51:
                             2598
                                                                                      2538
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                                                                                                                                                                                                    2448 ATGGCTGTGAACTGGCTCGGT---------GCAGGATGCCGGACAGAGGAC
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 78
                                                                                                                                                                       21 GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu 37
                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0.

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 4267 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/949,155
FILING DATE: <Unknown>
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WH
STREET: P.O. Box 4433
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Generation of Transgenic
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                                            ValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGlu 77
                            GTGCAGTCCCTGTCTGACCAAGTGCAGGAGGAGCTGCTCAGCACCAAGGTCACCCAGGAA
                                                                                     TGGCAGGGCAGCCCTGGGAGCAGGCCCTGGGCCGCTTCTGGGATTACCTGCGCTGG
                                                                                                  GAGCCGGGG-----
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ZIP: 77210-4433
                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Hibler, David W. REGISTRATION NUMBER: 41,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/819,964 FILING DATE: 28-Mar-2001
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48.94%
42.30%
58.04%
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Matches:
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Mismatches:
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RESULT 7
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                                                                                                                                                                                                                                           GENERAL INFORMATION:
                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                 APPLICANT: HORNES, Erik
APPLICANT: UHLEN, Mathias
TITLE OF INVENTION: CLONING
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                                                                                                                                                                                     NUMBER OF SEQUENCES:
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SOFTWARE: Patenti
JRRENT APPLICATION
                                                                                                                              STREET: 1800 Dia CITY: Alexandria
                                                                                                    COUNTRY:
                                                                                                                                                           ADDRESSEE:
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22313-0299
                                                                                                                 Virginia
                                                                                                                                           E: Foley & Lardner
1800 Diagonal Road
                 PatentIn Release #1.0,
       DATA
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Query Match:
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                                                                                                                                         Sequence 24,
Patent No. 60
                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 330 base pairs TYPE: NUCLEIC ACID
                                                                                 APPLICANT: Kvster, Hubert TITLE OF INVENTION: DNA D NUMBER OF SEQUENCES: 33
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REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 16787/168/DFBC
TELECOMMUNICATION INFORMATION:
                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: BENT, Stephen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.:
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  CITY: BC
STATE: M
COUNTRY:
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                                                        ADDRESSEE:
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                                             STREET:
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TELEFAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 19920
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeuThrProValAlaGluGlu
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                                                                                                                                                                                                                                                                                                                                           SerThrGluGluLeuArgValArgLeuAlaSerHisLeuArgLysLeuArgLysArgLeu 166
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Matches:
Conservative:
Mismatches:
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Best Local Similarity:
Query Match:
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                                                                                                                                         Sequence 24, Application US/09287141 Patent No. 6197498
GENERAL INFORMATION:
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REFERENCE/DOCKET NUMBER: SQ
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
                                                                                        APPLICANT: K ster, Hubert
TITLE OF INVENTION: DNA Diagnostics
NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/406/199
FILING DATE: March 17, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth A
REGISTRATION NUMBER: 35,430
                                                                          CORRESPONDENCE ADDRESS:
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LENGTH: 252 base pairs
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PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
        STREET:
CITY: I
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           109 AlaArgLeuSerLysGluLeuGlnÅlaAlaGlnAlaArgLeuGlyAlaAspMetGluAsp
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                                                          ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                       AspAlaAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla 188
                                                                                                                                                                                                                                                                                                                                                                                                    GluGluLeuArgValArgLeuAla$erHisLeuArgLysLeuArgLysArgLeuLeuArg
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                                       Heller Ehrman White & McAuliffe
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Indels:
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Best Local Similarity:
Query Match:
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                                                           Sequence 24, Application US/09431613 Patent No. 6221601
             GENERAL INFORMATION:
APPLICANT: K ster,
TITLE OF INVENTION:
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MOLECULE TYPE:
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LENGTH: 252 base pairs
TYPE: nucleic acid
NUMBER OF SEQUENCES:
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FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/6:
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
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TELEFAX: 619-587-5360
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REFERENCE/DOCKET NUMBER: 24736-2002D
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REGISTRATION NUMBER: 33,779
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OPERATING SYSTEM:
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(617)227-5941
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              Diagnostics Based
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Indels:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor

La Jolla

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Best Local Similarity:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TELEPHONE: 619-450-8400
TELEPAS: 619-587-5360
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: 08/40
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/617,256
FILING DATE: 18-MAR-1996
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24
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CLASSIFICATION:
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                                                    GluArgGlyLeu 192
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                                       GAGCGCGGCCTC
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Matches:
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Query Match:
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GENERAL INFORMATION:
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TELEFAX: 858-587-5360
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/617
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IIM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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LENGTH: 252 base pairs
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA Di
                                                                                                                                                                                                                                                                                                                                                                                                                                    No.:
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                                                                                                                             129 ValCysGlyArgLeuValGlnTyrArgGlyGluValGlnAlaMetLeuGlyGlnSerThr
                                                                                                                                                                                                                                                                            109 AlaArgLeuSerLysGluLeuGlnAlaAlaGlnAlaArgLeuGlyAlaAspMetGluAsp
                                                                181
                                                                                                                                                                                                62 GTGTGCGC-CGCCTGGTGCAGTACCGCGGCGAGGTGCAGGCCATGCTCGGCCAGAGCACC
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002J
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AspAlaAspAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla
                                                                GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGCCCGCGAGGGCCCC
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                     Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
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FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,1
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
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APPLICANT: Little, Daniel P
APPLICANT: Braun, Andreas
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                                                                            129 ValCysGlyArgLeuValGlnTyrArgGlyGluValGlnAlaMetLeuGlyGlnSerThr 148
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OPERATING SYSTEM: DOS
SOFTWARE: Patentin Release #1.0, Version
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002E
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                GluGluLeuArgValArgLeuAlaSerHisLeuArgLysLeuArgLysArgLeuLeuArg
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GAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGCAAGCTGCGTAAGCGGCTCCTCCGC
                                                             GTGTGCGC-CGCCTGGTGCAGTACCGCGGCGAGGTGCAGGCCATGCTCGGCCAGAGCACC
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Matches:
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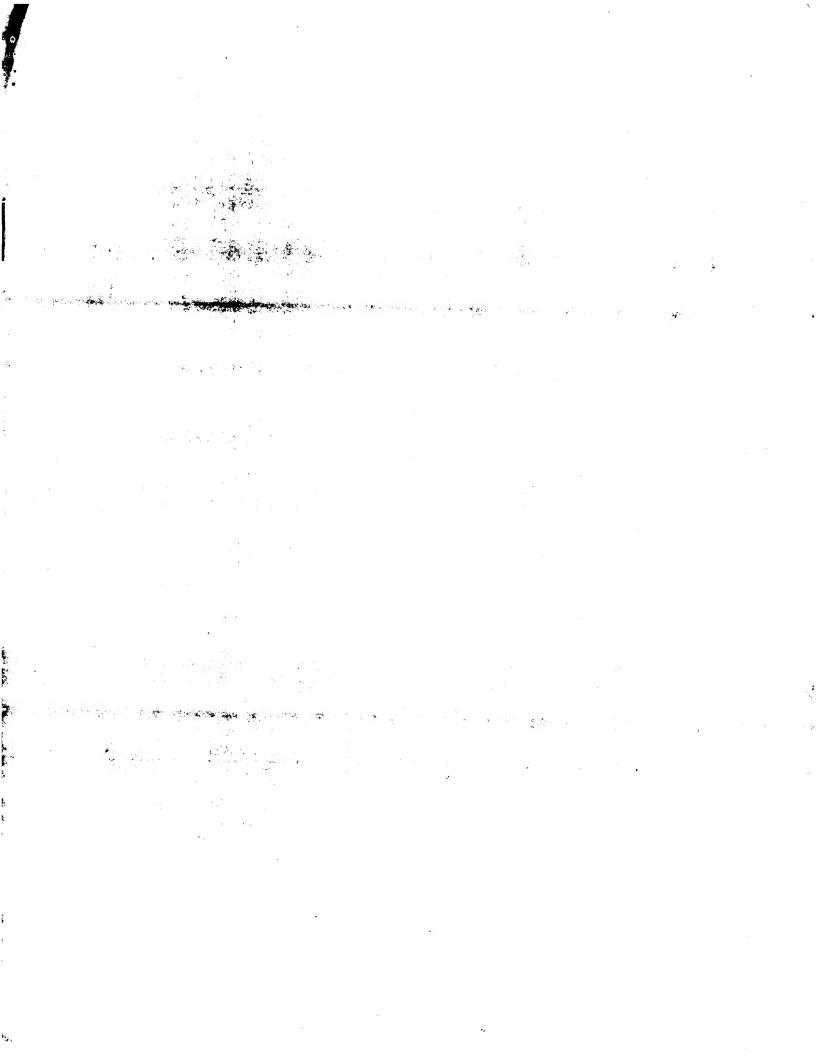
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                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/406,19
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 2473
TELECHONE: 619-450-8400
TELEPHONE: 619-450-8400
TELEFAX: 619-587-5360
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APPLICANT:
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MEDIUM TYPE: Diskette
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OPERATING SYSTEM:
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                                         Percent Similarity:
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TELEPHONE: (617)227-7400
TELEPHANE: (617)227-941
INFORMATION FOR SEQ ID NO: 24:
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APPLICATION NUMBER: 08/617
FILING DATE: 18-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/400
FILING DATE: 03-MAR-1995
                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 619-450-8400
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MEDIUM TYPE: Diskette
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ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS
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; TOPOLOGY: 1.
; MOLECULE TYPE:
US-09-287-681-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                TELEFAX: (617)227-594:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/617,256
FILING DATE: 18 MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/406,199
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM COmpatib
OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400
                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA Diagnostics Based on Mass Spectrometry NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              241
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                                                                                                                                                                                                                                                  NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,779
                                                 STRANDEDNESS: single
                                                                                                                                                                                    TELEFAX: 619-587-5360
                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 92037-9103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY:
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o. 6277573
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                                                                nucleic acid
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                                                                                                                                    (617)227-5941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM Compatible
                                                                                                                                                                                                     619-450-8400
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                                                                                                                                                                                                                                                                                                                                                                                                                                   US/09/287,681
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ER: 247
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Alignment Scores



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Minimum DB
Maximum DB
                                                                            Result
No.
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-QB-Published_App_ications_Na -QFMT=fastap_SUFFIX=p2n.rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX=bitosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -HR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -QUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=2000000000 -USER=US09827854 | @CCN_1_1_298 @runat_11032003_101612_27578
-NCPU=6 -ICPU=3 -NO_XLPXY -NO_MMAP +LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=*10 -YGAPOPXT=0.5 - DELOP=5 -QELEXT=7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 08
Maximum Match 10
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-MODEL=frame+_p2n.model
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                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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                                                                                                                                                                                                                                                                                                                                                                                                                         Published_Applications_NA: *
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-827-854-15_COPY_1_203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         501302 segs, 350932<mark>545 residues</mark>
                                                                                                                                                                                                         // Cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
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// Cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
// Cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-802-640-17
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2 US-10-044-090-454
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Sequence 129, App
Sequence 17, Appl
Sequence 8, Appli
Sequence 454, App
                                                                         Description
                                                                                                                                                                                                                                  US-09-870-759-129
; Sequence 129, Application US/09870759
; Patent No. USZ0020177551A1
; GENERAL INFORMATION:
APPLICANT: TERMAN, David S
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT OF NEOPLASTIC DISEASE
; FILE REFERENCE: 870759
; CURRENT APPLICATION NUMBER: US/09/870,759
; CURRENT FILING DATE: 2002-01-14
PRIOR APPLICATION NUMBER: US 60/208,128
; PRIOR APPLICATION NUMBER: US 60/208,128
; PRIOR FILING DATE: 2000-05-30
                                                                         ; NAME/KEY: CDS
; LOCATION: (61)..(1:
; OTHER INFORMATION:
US-09-870-759-129
 Percent Similarity:
                             Alignment Scores: Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      a a
                                                                                                                                                              NUMBER OF SEQ ID NOS: 166
SOFTWARE: PatentIn version 3.1
SEQ ID NO 129
LENGTH: 1156
TYPE: DNA
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US-09-827-854-9

US-09-827-854-11

US-09-954-456-760

US-09-880-107-2244

US-09-827-854-10

US-09-827-854-10
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              Length: Matches:
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                                                                                                                                                                                                                 Sequence 17, Application US/09802640 Publication No. US20030036057A1 GENERAL INFORMATION:
                                                                     SOFTWARE: FastSEQ
SEQ ID NO 17
LENGTH: 1156
                                                                                                                  APPLICANT: Braun, Andreas
APPLICANT: Bonsal Aruna
APPLICANT: Kleyn Patrick
TITLE OF INVENTION: GENES AND POLYMORPHISMS
TITLE OF INVENTION: CARDIOVASCULAR DISEASE
FILE REFERENCE: 24736-2048
CURRENT APPLICATION NUMBER: US/09/802,640
CURRENT FILING DATE: 2001-03-09
                                                                                                         NUMBER OF SEQ ID NOS:
           NAME/KEY: CDS
LOCATION: (61)
                                     FEATURE:
                                             ORGANISM: Homo sapien
                                                           TYPE: DNA
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Percent Similarity:
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DB:
GENERAL INFORMATION:
APPLICANT: Zannis, Vassilis
APPLICANT: Zannis, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of the compound of 
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SEQ ID NO 8
SEQ ID NO 8
; LENGTH: 1156
; TYPE: DNA
; ORGANISM: HOMO SAP
US-09-827-854-8
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Sequence 454, Application US/10044090
Patent No. US20020137081A1
GENERAL INFORMATION:
APPLICANT: Olga Bandman
TITLE OF INVENTION: GENES DIFFERENTIALLY EXPRESSED
FILE REFERENCE: PA-0028 US
CURRENT APPLICATION NUMBER: US/10/044,090
CURRENT FILING DATE: 2002-01-091
NUMBER OF SEQ ID NOS: 850
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RESULT 5
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Best Local Similarity:
Query Match:
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LENGTH: 1291
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TYPE: DNA
ORGANISM: Homo:
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OTHER INFORMATION:
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Best Local Similarity:
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TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/59,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
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             ProLeuVal
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; ORGANISM: HOMO
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Pred. No.:
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SOFTWARE: FastSEQ for
SEQ ID NO 9
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APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
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            GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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                                                   AAGCTGCGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGTGCCTGGCAGTGTAC
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Patent No. US/0020123093A1
GENERAL INFORMATION:
APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing hypertriglyceridemia
FILE REFERENCE: 07180/004003
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SEQ ID NO 11
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CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR EILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
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NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version
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TYPE: DN
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                                                CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
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AAGCTGTGTAAGCGGCTCCTCCGCGATGCCGATGACCTGCAGAAGCGCCTGGCAGTGTAC
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PRIOR APPLICATION NUMBER: US/60/234,052
PRIOR FILING DATE: 2001-09-18
PRIOR FILING DATE: 2000-09-18
PRIOR FILING DATE: 2000-09-24
PRIOR FILING DATE: 2000-09-20
PRIOR FILING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US/60/234,923
PRIOR APPLICATION NUMBER: US/60/234,923
PRIOR APPLICATION NUMBER: US/60/235,134
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR FILING DATE: 2000-09-25
PRIOR FILING DATE: 2000-09-26
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Best Local Similarity:
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; ORGANISM: Hom
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PRIOR APPLICATION NUMBER: US/60/235,638
PRIOR TILING DATE: 2000-09-26
PRIOR PELLOATION NUMBER: US/60/235,711
PRIOR APPLICATION NUMBER: US/60/235,720
PRIOR APPLICATION NUMBER: US/60/235,720
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US/60/235,840
PRIOR FILING DATE: 2000-09-27
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SEQ ID NO 760
LENGTH: 1157
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TITLE OF INVENTION: Sets
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CURRENT FILING DATE: 2001-09-18
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GAGCAAGCGTTGGAGACAGAGCCGGAGCCCGAGCTGCGCAGACCGAGTGGCAGAGC
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Best Local Similarity:
Query Match:
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CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Vockley
APPLICANT: Scherf,
APPLICANT: Gene Lo
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LENGTH: 1157
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TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5028-WO
                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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US-09-827-854-15_COPY_1_203 (1-203) x US-09-827-854-7 (1-1156)
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                                                    Percent Similarity:
Best Local Similarity:
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                                                                                                                                         ; ORGANISM: Homo US-09-827-854-7
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                                                                                                                                                                                             SEQ ID NO 7
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                                                                                                                                                                                                         COURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
NUMBER OF SEQ ID NOS: 19
NUMBER OF SEQ ID NOS: 19
NUMBER OF SEQ ID NOS: 19
NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                 APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Cholesterol levels without inducing hypertriglyceridemia
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
                                                                                                                                                                     TYPE: DNA
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MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal

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Alignment
Pred. No.:
                                                                                                            APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for TITLE OF INVENTION: Condesterol levels with File Reference: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
  Percent Similarity:
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Query Match:
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                                                                             TYPE: DNA
ORGANISM: Homo
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LENGTH: 1156
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; LENGTH: 41907
; TYPE: DNA
; ORGANISM: HOMO :
US-09-967-013-5
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US-09-967-013-5
Percent Similarity:
Best Local Similarity:
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                                                                                                                                                      SEQ ID NO 5
LENGTH: 41907
                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/206,613 PRIOR FILING DATE: 2000-05-23 NUMBER OF SEQ ID NOS: 91
                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/967, CURRENT FILING DATE: 2000-10-25
                                                                                                                                                                                                                                                                                          APPLICANT: Stanton, Jr. Vincent P. TITLE OF INVENTION: METHOD FOR GENETIC TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: 11926-022001
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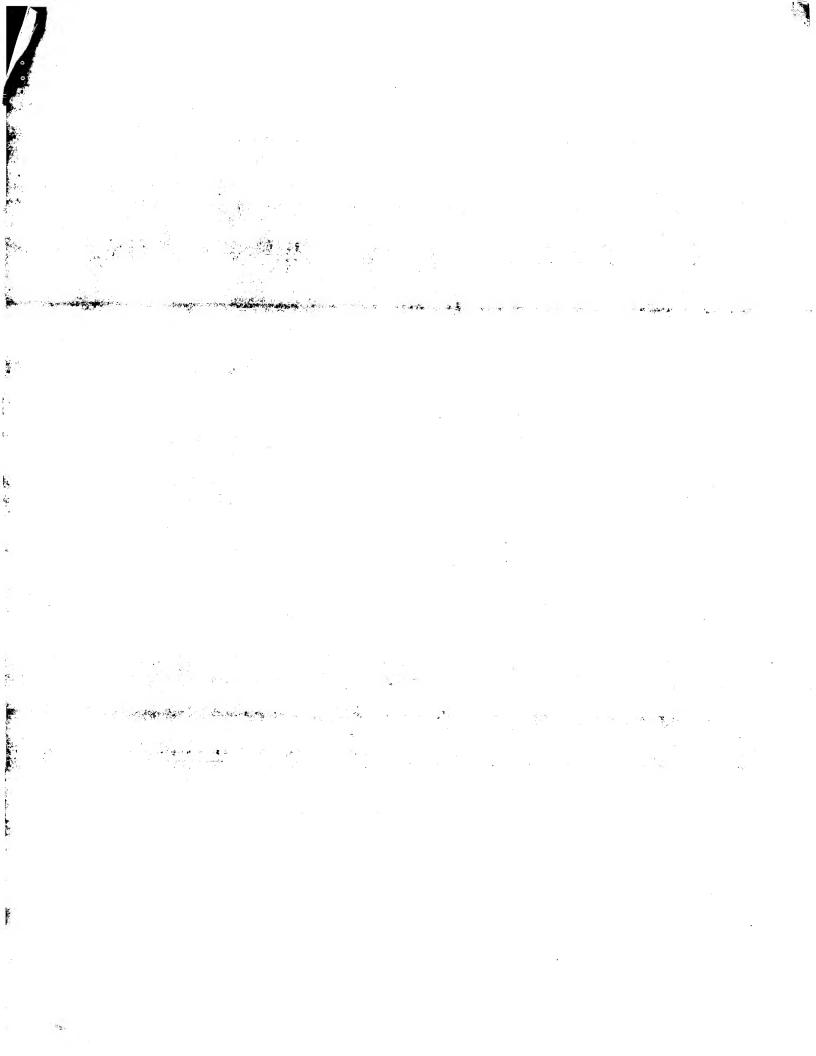
21 15 21	CAGGCCCGGCTGGGCGCGGACATGGAGGACGTGYGCGGCCGCCTGGTGCAGTACCGCGG 2 GluValGlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHi 1	Db Db
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	20556 GGCGGCTATACCTCCCCAGGTCCAGGTTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGGG	Db
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	72 SerGlnValThrGlnGluLeu	Qy Db
	52 TrpAspTyrLeuArgTrpValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSer 	Qу
	32 LeuArgGlnGlnThrGluTrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPhe 	Оy
	12 PheLeuAlaGlyCysGlnAlaLysValGluGlnAlaValGluThrGluProGluProGlu	Qy Db
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SEQ ID NO 374
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TYPE: DNA
ORGANISM: Homo sapiens
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LOCATION: (1)...(478)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/964,824A CURRENT FILING DATE: 2001-09-27 PRIOR APPLICATION NUMBER: US/60/236,033 PRIOR FILING DATE: 2000-09-28 PRIOR APPLICATION NUMBER: US/60/236,032 PRIOR APPLICATION NUMBER: US/60/236,032 PRIOR FILING DATE: 2000-09-28
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PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
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Best Local Similarity:
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APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-06-19
PRIOR FILING DATE: 2000-06-19
PRIOR FILING DATE: 2000-06-19
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; LOCATION: (1)...(478)
; OTHER INFORMATION: n =
US-09-880-107-2491
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2491
LENGTH: 478
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APPLICANT: Vockle:
APPLICANT: Scherf,
APPLICANT: Gene Lo
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OTHER INFORMATION: Genbank Accession No.
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; LCCATION: (430)
; OTHER INFORMATION: unsure at all n locations
; OTHER INFORMATION: Clone ID: 20-LIB34-078-Q1-E1-E7
US-09-960-352-4726
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DB:
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APPLICANT: Tao, Nenghing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECU
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001.09-24
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SEQ ID NO 4726
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ORGANISM: BOS 1
FEATURE:
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                                                                 CTGATGGAGGAGCCATGAAGGAGGTGAAGGCCTACAAGGAGGAGCTGGAGGGACAGCTA
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Search completed: March 15, 2003, 03:04:06 Job time : 52.1851 secs



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Command line parameters:
-MODEL-frame+_p2n.model -DEV=xlp
-Q-/cgn2__/USPT0_spool/US09827854/runat_11032003_101610_27495/app_query.fasta_1.3576
-Q-/cgn2__/USPT0_spool/US09827854/runat_11032003_101610_27495/app_query.fasta_1.3576
-DE-EST -QFMT=fastap -SUFFIX=p2n.rsft -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DCCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTEMT=ptc -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=20000000000
-USER=US09827854_@CCN_1_1_8826_@runat_11032003_101610_27495 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOF=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Maximum DB seq length: 200000000
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## SUMMARIES

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## ALIGNMENTS

TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BI599793	RESULT 1	
National Institutes of Health, Mammalian Gene Collection (MGC)	NIH-MGC http://mgc.nci.nih.gov/.	1 (bases 1 to 690)	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	human.	EST.	BI599793.1 GI:15492732	BI599793	mRNA sequence.	603245228F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5287710 5',	BI599793 690 bp mRNA linear EST 07-SEP-2001			

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                                            CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCCGCCTGGTGCAGTACCGCGGCGAGGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), (
TOshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLAM11727 row: g column: 07
High quality sequence stop: 686.
Location/Qualifiers
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Robert Strausberg,
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/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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US-09-827-854-15_COPY_1_203 (1-203) x BG769455 (1-699)
                                                                                                             Percent Similarity:
Best Local Similarity:
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 699)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
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Tissue Procurement: ATCC/DCTD/DTP
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Contact: Robert Strausberg, Ph.D.
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                    Plate: LLAM10676 row: 1 column: High quality sequence stop: 748. Location/Qualifiers
                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
Toshiyuki and Piero Caminici (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.
cDNA Library Arrayed by:
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602672086F1 NIH_MGC_96
                                                             DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution i found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
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/organism="Homo
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/clone="IMAGE:4794789"
/clone_lib="NIH_MGC_96"
/tissue_type="hypothalamus"
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                                                LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Toshlyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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National Institutes of Health, Mammalian
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Tissue Procurement: Miklos Palkovits, M.F.
cDNA Library Preparation: Michael J. Brc
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E.
                                                                                                                                                                                                                                                                                                                             High quality sequence stop: Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                 DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11768 row: d column: 21
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collections, 1000
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/note-"Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

230 c 292 g 97 t
                                                                                                                                                                                                                               /organism="Homo sapiens"
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/clone="IMAGE:5303396"
/clone_lib="NIH_MGC_96"
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       Homo sapiens
Eukaryota; Metazoa; Chordata; Gra
Mammalia; Eutheria; Primates; Cat
1 (bases 1 to 757)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Ma
Unpublished (1999)
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Tissue Procurement: DCTD/DTP
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/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: xhoI; Site_2:
/note="Organ: skin; Vector: pOTB7; Site_1: xhoI; Site_2:
ECORI; CDNA made by oligo-dT priming Directionally clone
into EcoRI/XhoI sites using the following 5: adaptor:
GGCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California)
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                                                                                                                                                                                                                                                                  BASE COUNT
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                              ATGAAGGTTCTGTGGGCTTGCTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTG
                                                                                                                                                      Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA sequence.
BI551811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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603192716F1 NIH_MGC_95
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                                                                                                                                                                                                                                                               /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: Sal-Xhol (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.5 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

a 235 c 311 g 103 t
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/clone="IMAGE:5264115"
/clone_lib="NIH_MGC_95"
                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="hippocampus"
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61 LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla 80
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                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI),
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM10677 row: c column: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 798)
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BG708414.1 GI:13985733
EST.
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                                                                                            Plate: LLAM10677 row: c column: High quality sequence stop: 778.
                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999
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National Institutes of Health, Mammalian Gene Collection (MGC)
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/organism="Homo sapiens"
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                                                                        location/Qualifiers
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/clone="IMAGE:4794955" /clone\_lib="NIH\_MGC\_96"

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RESULT 9
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          BM042094
603615713F1 NIH_MGC_112
mRNA sequence.
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/lab_host="DH10B"
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Best Local Similarity:
Query Match:
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                                                                                               LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluGlnLeu 100
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                                                                      CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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                                                                                                                                                                                                                                                                   GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTTGGGATTACCTGCGCTGGGTGCAGACA
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Tissue Procurement: DCTD/DTP
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Contact: Robert Strausberg, Ph.D.
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/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
ECCRI; cDNA made by ollgo-dT priming. Directionally cloned
into EcCRIXhoI sites using the following 5' adaptor:
GGCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
NIH_MGC Library."

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/db_xref="taxon:9606"
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National Institutes of Health, Mammalian
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                     cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1612 row: e column: 20 High quality sequence stop: 813. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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1 (bases 1 to 817)
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                                                                                              /clone_lib="NIH_MGC_40"
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/lab_nost="DH1DB (phage-resistant)"
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/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
/site_2: EcoRI; cDNA made by oligo-dT priming,
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Supersoript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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/clone="IMAGE:4760827"
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1 (bases 1 to 845)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                BG829472.1
EST.
                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
                                                                                                                                                 Unpublished (1999)
Contact: Robert Strausberg,
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                 CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M. Sequencing by: Incyte Genomics, Inc. Consortium (Clone distribution: MGC clone distribution information)
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Plate: LLCM1790 row:
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH20B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRIXhoI sites using the
following 5' adaptor: GGCAGGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Ling
Hong in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NH_MGC Library. | "
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/clone="IMAGE:4899112"
/clone_lib="NIH_MGC_42"
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TITLE
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), SP
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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Plate: LLAM11798 row: a column:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
/clone="IMAGE:5314844"
/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1999)
Contact: Robert Strausberg, F
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 965)
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5', mRNA sequence.
BQ677266
                                                                                                                                                                                                                                                                                 CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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US-09-827-854-15_COPY_1_203 (1-203)
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EST.
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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            ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal 140
                                                                        ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
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CGGCTGGGCGCGGACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGATGTG
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Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov http://image.lnnl.gov plate: LLAM11662 row: g column: 01
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Query

Match:

US-09-827-854-15\_COPY\_1\_203 (1-203)

x BI458355 (1-802)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tissue Procurement: Miklos Palkovits, M.D., Ph.D. cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh Toshiyuki and Piero Caroinci (RIKEN) cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 
Location/Qualifiers
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http://image.llnl.gov
Plate: LLAMI1702 row: j column:
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
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                                                                                                                                                                             143
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                                                                                                                                                                         /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site-1: BamHI; Site-2: SalI-Xhol (gtcg: pBluescript KS+); Site-1: BamHI; Site-2: SalI-Xhol (gtcg: pBluescript KS+); Site-1: BamHI; Site-2: Skb and contailized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."

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                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
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Search completed: March 14, 2003, 20:13:59 Job time: 829.891 secs

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-DB=GenEmbl -QFWT=fastap -SUFFIX=p2n.rge -MINNATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END--1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN=15 -MODE-LOCAL
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-WARN_TIMEOUT=30 -THREADS=1 -XGAPOE=10 -XGAPEXT=7.5 -FGAPOP=6 -FGAPEXT=7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

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                                            CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
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MITSUBISHI CHEM IND LTD

OS human

PN JP 1985118189-A/1

PD 25-JUN-1985

PFT CDS

PN JP 1985118189-A/1

PD 25-JUN-1983

PN JP 1983224

PI TERANISHI YUTAKA, TAKA

KINURA MASAKO,

PI IKEDA YASUKO

PC C12N15/OD,CO77H21/O4//C

CC strandedness: Double;

CC topology: Linear;

CC hypothetical: No;

CC anti-sense: No;

CC anti-sense: No;

CC atti-sense: No;

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Teranishi,Y., Takamatsu,N., MaDNA FRAGMENT
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29-NOV-1983 JP 1983224980
TERANISHI YUTAKA, TAKAMATSU
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strandedness: Double;
topology: Linear;
hypothetical: No;
anti-sense: No;
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/db_xref="taxon:9606"
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    Location/Qualifiers
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Matches:
Conservative:
Mismatches:
Indels:
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1986096997-A/1.

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unidentified
unclassified.
1 (bases 1 to 1110)
1 (bases 1 to 1110)
2 PACOUCTION OF HUMAN APOLIPOPROTEIN E-LIVER AND LTD
DOWNENT OS HUMAN (HOMO SAPIENS)
PN JP 1986096997-A/1
PD 15-MAY-1986
PF 16-OCT-1984 JP 1984216987
PI TERANISHI YUTAKA, MATSU'
PC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C12N15/00, FC (C
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15-MAY-1986
15-MAY-1986
16-OCT-1984 JP 1984216987
TERANISHI YUTAKA, MATSUI YASUSHI, IKEDA YASUKO, KIMURA MASAKO C12P21/00, A61K35/74, A61K37/04, C12N15/00, (C12P21/00, C12R1:19), (C12N15/00,
                        CDS
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                                                                                                                                                           tissue_type=Liver;
    Location/Qualifiers
                                       /product='human
peptide'
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                        69.
/product='human
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Wallis,S.C., Rogne,S., Gill,L., M
Williamson,R. and Humphries,S.
The isolation of cDNA clones for
detection of apoE RNA in hepatic
EMBO J. 2 (12), 2369-2373 (1983)
                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Mammalla; Eutheria; Primates; Catarrhid 1 (bases 355 to 1156)
Breslow, J.L., McPherson, J., Nussbaum, A.L Lofquist-Kahl, F., Karathanasis, S.K. and
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/organism="Homo sapiens" /db_xref="taxon:9606" /map="19q13.2" /tissue_type="liver and blood" 1. 1156 /gene="APOE"	A- see [J] re an Ap PN Ap CO CO CO CO CO L. CO CO CO CO CO CO CO CO CO CO CO CO CO	variant Journal J. Clin. Invest. 83 (4), 1095-1101 (1989)  MEDLINE 89198059  PUBMED 2539388  [1] corrected in [J. Biol. Chem. 258, 11422-11422 (1983)]. [J. Biol. Chem. 258, 11422-11422 (1983)]. [J. Chem. 258, 1	REFERENCE 3 (bases 1 to 1156) AUTHORS Zannis,V.I. McPherson,J., Goldberger,G., Karathanasis,S.K. and Breslow,J.I. Synthesis, intracellular processing, and signal peptide of human JOURNAL J. Biol. Chem. 259 (9), 5495-5499 (1984)  MEDLINE 6325438  REFERENCE 4 (bases 88 to 1156) AUTHORS 743(7.J.M. Elshourbagy,N.A., Chang,D.J., Mahley,R.W. and TITLE of a new variant JOURNAL J. Biol. Chem. 259 (10), 6498-6504 (1984)  MEDLINE 64212473  MEDLINE 6212473  REFERENCE 5 (bases 577 to 624) AUTHORS 6111,L.L., Peoples,O.P., Pearston,D.H., Robertson,F.W., Humphries,S.E., Cumming,A.M. and Hardman,N.  TITLE JOURNAL Biochem. Biophys. Res. Commun. 130 (3), 1261-1266 (1985)  REFERENCE 6 (sites) REFERENCE 7 (sites) REFERENCE 8 (1985) REFERENCE 8 (1985) REFERENCE 8 (1985) REFERENCE 9 (1984)  MEDLINE 9292507 REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 6 (1985) REFERENCE 7 (1984) REFERENCE 7 (1984) REFERENCE 8 (1985) REFERENCE 8 (1985) REFERENCE 9 (1984) REFERENCE 9 (19
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CIREII. 230, 11422 <sup>-</sup> 11422 (1903)],[3],	oproteinemia" oproteinemia"	Chem. 258, 11422-11422 (1983)],[3]"	611014 /gene="APOE" /codon_start=1 /product="preapolipoprotein E" /protein_id="AAB59446.1" /protein_id="AAB5946.1" /db_xref="GI:178851" /db_xref="GI:178851" /db_xref="GB:GO0-119-691" /db_xref="GB:GO0-119-691" /db_xref="GB:GO0-119-691" /db_xref="GB:GO0-119-691" /db_xref="GB:GO0-119-691" /db_xref="GB:GO0-119-691" /db_xref="GB:GO0-119-691" /protein_id="AKVLMAALLVTFLAGCQAKVEQAVETEPEPELRQOTEWQSGORW /translation="MKVLMAALLVTFLAGCQAKVEQAVETEMELKAYKSELEEQLTP /EARACAMGERLARMEMASLVEKUQAAVETSAALMDETMKELKAYKSELEEQLTP VARETRARLSKELQAQAARLGAMMEDVGERLUQVERGEVQAMIGQSTEELKAYKSELEEQLTP VARETRARLSKELQAQAARLGAMMEDVGERULGAVAGSLAG QPLQERAQAMGERLARMEMMGSRTRDRLDEVKEQVAEVRAKLEEQAQQIRLQAEAFQ ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH" /gene="APOE" /product="apolipoprotein E mature peptide" /gene="APOE" /product="apolipoprotein E mature peptide" /gene="APOE" /citation=[2] /replace="c" /gene="apoE" /gen

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Tissue Procurement: ATCC
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VAEETRAFLSKELQAAQARLGADBEDVCGRLVQYRGEVQAMLGQSTEELKVRLASHLR
KIRKRLLRDADDLQKRLAYYQAGAREGAERGLSAIRERLGPLVEQGRVRAATVCSLAG
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/lab_host="DH10B-R"
/note="Vector: pOTB7"
46. 999
                          ARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDNH" 366 c 425 g 147 t
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/db_xref="LocusID:348"
/db_xref="Laxon:9606"
/clone="MGC:1571 IMAGE:3355712"
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                                                                                    V DNA coding human apolipoprotein E08423 E08423 E08423 E08423 GI:2176540 JP 1994315392-A/1. Homo sapiens Homo sapiens Eukaryota; Metazoa; Chordata; C
                                                                                                                           SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
                                                                                                                                                                                                              CAGGCCGGGGCCCGAGGCGCGGCCTCAGCGCCATCCGCGAGCGCCTGGGG
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Craniata; Vertebrata;
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                                       GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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MORIMOTO, H. and Teranishi, Y.
METHOD FOR PRODUCING APOLIPOPROTEIN
Patent: JP 1994315392-A 1 15-NOV-1994;
MITSUBISHI KASEI CORP
OS HOMO Sapiens (human)
PN JP 1994315392-A/1
PD 15-NOV-1994
PF 11-JUN-1985 JP 1994015433
PI MORIMOTO HIRONORI, TERANISHI YUTAKA
PC C12P21/02,COYK13/00,C12N5/10,C12N15/
PC C12R1:91),
PC C12R1:91),
PC C12R1:91);
PC C12R1:91);
PC C12R1:91);
PC C12R1:91);
PC C12R1:91);
PC topology: Linear;
FH Key
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FT sig_peptide
FT mat_peptide
FT //product='human apol
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LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                                            ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                                ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
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                                                                               CGGCTGGGCGCGCACATGGAGGACGTGTGCGGCCGCCTGGTGCAGTACCGCGGCGAGGTG
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MORIMOTO HIRONORI, TERANISHI YUTAKA
C12P21/02,C07K13/00,C12N5/10,C12N15/12,C12N15/18,(C12P21/02,
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21
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1 (bases 1 to 1156)
5 Fujita, S., Hamanaka, H., Fukui, Y. and Yokoyama, M. Apo E humanized mammal patent: JP 2001017028-A 1 23-JAN-2001; Patent: JP 2001017028-A 1 23-JAN-2001; Patent: JP 2001017028-A/1
PN JP 2001017028-A/1
PN JP 2001017028-A/1
PD 23-JAN-2001
PF 28-APR-2000 JP 2000128919
PR SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINE PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINE
                             GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeuGly
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BD004277.1 GI:1863:
JP 2001017028-A/1.
                                                                                                                                                                                                                                                                                                                                                                                                                          PI SHINOBU FUJITA, HIROKI HAMANAKA, YUKO FUKUI, MINESUKE A01K67/027, A61K45/00, A61P25/28, A61P43/00, C12N5/10, PC C12N15/09//C07K14/775, PC (C12N5/10.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1.C12P1
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Location/Qualifiers
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/db_xref="taxon:9606"
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Matches:
Conservative:
Mismatches:
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                                                                   Avalon
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LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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                                                                                                                                                                                                                                                                                                                                                                  AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCGCCAAG
                                                                                                                                                                                                                                                                                                                                                                              SerargThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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Sequence
AX333278
AX333278.
                                                                                                                                              Homo sapiens
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                         Young, P.E., Augustus, M., Carter, K.C., Horrigan, S., Soppet, D.R. and Weaver, Z
                                                                                sets
                           : WO 0194629-A 3787 13-DEC-2001;
Pharmaceuticals (US)
Location/Qualifiers
1. .1157
                                                                                             gene
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/organism="Homo sapie
/db_xref="taxon:9606"
                                                                                             , Soppet, D.R. a determination
                                                                                                                                                                                                                     GI:18123912
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                                                                                                                                                                                                                                              Patent
                                                                                                                                                   Chordata; Craniata; Vertebrata; Primates; Catarrhini; Hominidae;
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                                                                                             and
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WO0194629
                                                                                             therapeutic
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AX409597 LOCUS DEFINITION	Db 8								Ov 1			Qy Db 1	Qy Db 1	Оy	US-09-827-854-15_COPY_1_277 (1-277	Alignment Pred. No.: Score: Percent Si Best Local Query Matc DB:	BASE COUNT ORIGIN
ž	, v =		0-1	0-						81 Lo 302 Ci		41 GI     182 G	21 GJ 	1 Me 62 As	27-85	nment Sc. No.: e: Ent Simi Local S. y Match:	TNUC
AX409597 Sequence	LeuGluGluGLnAlaGlnGlnTleArgLeuGlnAlaGluAlapheGlnAla	Technology   Tec	LeuG.InGluArgAlaGInAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly	ProLeuValGLuGLuGLyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro	GINALGGIYALBARGGLUGIYALBGIUARGGIYLEUSERALBILBARGGIUARGLEUGIY	LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr				TeumetAspGluThrMetLysGluTeuLysAlaTyrLysSergIuLeuGluGluGluGluCluLysAlaTyrLysSergIuLeuGluGluGluGluCluLysAlaTyrLysSergIuLeuGluGluGluCluLysAlaTyrLysSergIuLeuGluGluGluCluLysAlaTyrLysSergIuLeuGluGluCluCluCluCluCluCluCluCluCluCluCluCluCl	LeuSerGlucInValcInGlucIuLeuLeuSerSerGlnValThrGlnGluLeuArgAla	GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr 	GluGlnAlaValGluThrGluProGluProGluLeuArgGlnGlnThrGluTrpGlnSer 	MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal 	54-15	ores lari imil	
9597 ence	GLUGI	ACCCC	GLUAR         GAGCG	ValGI	GEGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	ArgLy	ATGCT	Met L	GTAGO	AspGJ        GACGP	GAGC <i>I</i> 	ArgTı        CGCTC	Alava        GCGG1	Valle        GTTCT	COPY	ty: .arity	212 a
2244	nAlac       GGCC	CGACO	gAlac       GGCC	HIII ACAG	aArgo	'sArgi       GCGG	1111	IIII	GGAG	LuThri       GACC	LnVal	rpGlu       3GGAA	liglu IIIII Iggag	euTrp.	[_1_2		
from	31nG1        CAGCA	CGCCT	31nA1	GlyAr        GCCG	GAGGG	CTCCT	CAGAG	ATGGA	HATEL	MetLy       ATGAA	GlnG1       CAGGA	LeuAl       CTGGC	ThrG1	AlaAl       GCTGC	77 (1	7.62e-7; 1365.00 99.28% 99.28% 99.38%	370 c
Pate	nIleA       GATAC	HIIII	aTrpG	gValA           CGTGC	yAlac           CGCCG	uArgA        CCGCG	CACCG	GGACG	GCGGG	sGluI        GGAGT	uGluI        GGAGC	aLeuc        ACTGG	uProc        GCCGC	aLeuI       GTTGC	-277)	72	
	rgLeu 	AGGTG	lyGlu             GCGAG	rgAla                GGGCC	LuArg	SPAla	AGGAG	TGTGC	CACGG	euLys	euLeu TGCTC	lyArg        GTCGC	luPro           AGCCC	euVal        TGGTC	×	0 M Z O Z H	26 g
157 bp [ W00229103.	GlnAl         CAGGC	AAGGA	ArgLe	ALATI         GCCAC	GGLYL6	ASPAS        GATG <i>I</i>	CTGC	I I I I I	CTGTC	AlaTy	Serse	Phen 	GluLe        GAGCT	ThrP	AX333278	Length: Matches: Conservative Mismatches: Indels: Gaps:	149
DNA	aGlu <i>t</i> 	GCAGG	uArg <i>i</i> 	TGTGC	uSer/        CAGC	CCTGC	HILL	CCTGC	CAAGO	TLYS	erGln\       CCAA	PASP	euArgo        GCGCO	neLeu/   - - -  CCTG	78 (1	n: es: crvativ	t 61
	laPhe      CCTT	TGGC	llaAr       CGCG	llyse:       GCTC	Lail   CCAT	lnLy:       AGAA	CCTC	TGCA	BAGCT	SerGL	/alTh	TYTLE       TACCT	BlnGl: BAGCA	AlaGl        GCAGG	(1-1157)		
linear	GlnA        CAGG	GAGG	MetG.	CLeuA	Arge         CGCG	ArgL        cccc	CGCCT	TACC	CAGA	ACTGG	rGlnG        CAAG	ArgT.       3CGCT	Thre	YCYSG       ATGCC	Ŭ	1157 275 0 2 0	
	la 277    CC 892	recec	luGlu       AGGAG	laGly        CGGC	luArg          AGCGC	euAla        rGGCA	CCCAC	GCGGC	00000	luGlu       AGGAA	luLeu       AACTG	rpVal       GGGTG	luTrp       AGTGG	lnAla       AGGCC			
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14-JUN-2002			. , -	, -	, ,					, ,				<b>-</b>			
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euVal        TGGTG	laGly         CCGGG	euArc        TGCGT	LaMet	euGl)         TGGGC	roVa]        CGGT#	letAsi          TGGAC	erGlu	lnare         AGCGC	lnala         AAGCG	ysval        AGGTT	15_cc	rity:	212		GENE LOGIC	human. Homo sapier Eukaryota; Mammalia; I 1 Alvares,C., Gene expres	AX409597 AX409597.1
ProLeuValGluGl           CCCCTGGTGGAACA	GlnAlaGlyAlaAr            CAGGCCGGGGCCCG	LysLeuArgLysAr             AAGCTGCGTAAGCG	GlnAlaMetLeuGl             CAGGCCATGCTCGG	ArgLeuGlyAlaAs             CGGCTGGGCGCGGA	ThrProValAlaG1	LeuMetAspGluTh	LeuSerGluGlnVa	GlyGlnArgTrpGl             GGCCAGCGCTGGGA	GluGlnAlaValGl               GAGCAAGCGGTGGA	MetLysValLeuTr           ATGAAGGTTCTGTO	Match: -827-854-15_COPY_1_	tv:	2 a / ".	1 1 0 1 1 0 1 1 0 1 1 0 1 1 0 1 1 1 0 1 1 1 1 0 1	TOGIC :	sapiens ryota; N alia; Et express	97 97 . 1

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Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
bb_xref="taxon:9606"
note="EMBL/GenBank Accession No. M12529"
370 c 426 g 149 t
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                                   7alGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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                                                                                                                                                                                                                               CCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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                       GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgfTrpValGlnThr
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Attie,A.D., Gretch,D.G., Sturley,S.L.
Production of recombinant proteins in
Patent: US 5472858-A 1 05-DEC-1995;
Location/Qualifiers
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/note="precursor"
/codon_start=1
/codon_start=1
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VAEETRAFLSKELQTAQARLGADAREGAERGLSAIRERLGPLVEQGRVWAATVGSLAG
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/note="apoE mRNA;
62. .1015
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<1. .1157
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1. .1157
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/sex="male"
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/map="19q13.2"
/clone="pHAE[112,178,813
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                                                                                                                                     /gene="APOE"
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**Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

**Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.**

**AUTHORS Fujita, S., Hamanaka, H., Fukui, Y. and Yokoyama, M.**

**TITLE Fujita, S., Hamanaka, H., Fukui, Y. and Yokoyama, M.**

**TITLE Apo E humanized mammal Patent: JP 2001017028-A 3 23-JAN-2001;

**MATENTICHEMICAL CORP OS Homo sapiens (human)

**PN JP 2001017028-A/3

**PD 23-JAN-2001

**PR 28-APR-2000**

**PR PR PR PAT 31-JAN-2

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                                                                                                                                MFAPOE 1178 bp
Monkey mRNA for apolipoprotein E
X13887
X13887.1 GI:38054
apolipoprotein; apolipoprotein E
Macaca fascicularis.
                                                                                                                                                                                                                                                                                                            CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
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                                                              Mammalia; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Cercopithecinae; Macaca.

1 (bases 1 to 1178)

Marotti,K.R.
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 Direct Submission
Submitted (16-DEC-1988) Marotti K.R., The
301 Henrietta Street, Kalamazoo, MI 49008
2 (bases 1 to 1178)
Marotti,K.R., Whitted,B.E., Castle,C.K., F
Melchlor,G.W.
Nucleotide sequence of the cynomolgus monk
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apolipoprotein F
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                                               ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
 CAGGCCATGCTGGGCCAGAGTACCGAGGAGCTGCGGGCGCCTCGCCTCCCACCTGCGC
                                      CGGCTGGGTGCCGACATGGAGGACGTGCGCAGCCGCCTGGTGCAGTACCGCAGCGAGGTG
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137. .103
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1154. .1159
/note="polyA signal"
373 c 426 g 162
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VAEETRARLSKELQAAQARLGADREDVRSRLVQYRSEVAAMLGQSTEELRARLASHLS
KLRKRLLRDADDLQKRLAVYQAGAREGAERGVSQIRERLGPLVEQGRVRAATVGSLAS
QPLQERAQALGERLRARMEEMGSRTRDRLDEVKEQVAEVRAKLEEQAQQISLQAEAFQ
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/clone="1"
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/protein_id="CAA32092.1"
/db_xref="GI:38055"
/db_xref="SWISS-PROT:P10517"
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/clone_lib="lambda g
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1 (bases 1 to 5491)
Nickerson,D.A., Taylor,S.L., Fullerton,S.M., Weiss,K.M., Clark,A.G., Stengard,J.H., Salomaa,V., Boerwinkle,E. and Sing,C.F Sequence diversity and large-scale typing of SNPs in the human apolipoprotein E gene Genome Res. 10 (10), 1532-1545 (2000)
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University of Washington, Bos
Location/Qualifiers
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Nickerson, D.A.
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                                                               /replace="<320. .>33
                                                                                                                                                                                                                                                                                            /map="19q13.2"
<3. .>108
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                    /rpt_type=tandem <340. .>637
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/note="putative"
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/note="putative"
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/db_xref="taxon:9606"
/chromosome="19"
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Qy 72 SerGlnValThrGlnGluLeu 78
Qy 52 TrpAspTyrLeuArgTrpValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSer 71
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US-09-827-854-15_COPY_1_277 (1-277) x AF261279 (1-5491)
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"g" c.
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78		78	Оу
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Search completed: March 14, 2003, 17:32:18 Job time: 2022.1 secs

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score greater than or equal to and is derived by analysis of Pred. No. is the number of results predicted by chance to have ster than or equal to the score of the result being prived by analysis of the total score distribution. printed, Ŋ

#### SUMMARIES

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37.4 36.7 35.3 30.5	0 5 5 5 5 7 7 9 9	73.1 73.1 73.0 71.6 66.9 64.1 63.5	00000000000000000000000000000000000000	Query Match
330 405 345 260	07770808	5617 6026 1126 660 936 936 597	111 111 111 111 111 111 111 111 111 11	Length
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Human apolipoprote Human secreted pro Human EST-derlved Human apolipoprote	cDNA encodi lone CP147. cancer assoc cDNA for no #2492 used t adenocarcin d cancer re	alpha album Lide Lide hum l hum x2 pr x2 pr x2 pr x2 pr x2 pr	man apolipopro an apolipoprotein E man ApoE3 codi man ApoE3 codi man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man ApoE4 codi man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man apolipopro man ApoE genom man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno man ApoE geno	Description

### ALIGNMENTS

RESULT 1 AAD26035 FH XXX AAD26035; AAD26035 standard; cDNA; Human apolipoprotein E (APOE) cDNA. 26-MAR-2002 (first entry) 954

Human; antilipaemic; neuroprotective; nootropic; genetic variant; APOE; apolipoprotein E; haplotyping; familial dysbetalipoproteinaemia; therapy; genotyping; type III hyperlipoproteinaemia; Alzheimer's disease; SNP; atherosclerosis; single nucleotide polymorphism; chromosome 19q13.2; ss.

Homo sapiens.

Location/Qualifiers

кеу

(1-954)

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Alignment
Pred. No.:
Score:
                                                                                                    Percent Similarity:
Best Local Similari
               Query Match:
                                                                                                                       The patent discloses novel genetic variants of human apolipoprotein CC E (APOE) gene. The invention also relates to compositions and methods CC for haplotyping and/or genotyping the APOE gene. The haplotyping CC methods of the invention are useful for improving the efficacy and CC reliability of several steps in the discovery and development of CC drings for treating diseases associated with APOE activity, e.g. CC familial dysbetalipoproteinaemia, type III hyperlipoproteinaemia, CC atherosclerosis, and Alzheimer's disease. They are useful to validate APOE as a candidate agent for treating a specific condition or disease predicted to be associated with APOE activity and in the design of C clinical trials of candidate drugs for treating a specific condition CC or disease predicted to be associated with APOE activity. Genotyping C or haplotyping methods are useful to screen for compounds targeting APOE to treat a specific condition or disease associated with APOE activity. The present sequence is a CDNA encoding human APOE protein.

CAPOE gene is located on chromosome 19q13.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Choi
                                                           No . :
                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDS
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                           Similarity:
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Location/Qualifiers
                                    hyperlipidemia;
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Best Local Similarity:
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LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                               CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
                                           GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                      ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                                ThrProValAlaGluGluThrArgAlaArgLeuSerLySGluLeuGlnAlaAlaGlnAla
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                                                                           CGGCTGGGCGCCGACATGGAGGACGTGTGCGGCCGCCCTGGTGCAGTACCGCGGCGAGGTG
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                                                                                                                                                               CTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAACTG
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The invention relates to methods for diagnosing and prognosing over the tumours in an individual via the detection and measurement of the expression of ovarian tumour marker genes (ABA83181-ABA83122, ABA83180, ABA83182 and ABA83184) or segments thereof (ABA83123-ABA83169, ABA83179, ABA83181 and ABA83183). The methods of the invention are useful for ABA83181 and ABA83183) in a patient, for identifying an individual

Detecting and identifying ovarian tumor, id developing ovarian cancer, and determining cancer treatment, by measuring expression l

identifying increased risk for ng effectiveness of ovarian n level of ovarian tumor marker

Claim

23;

Page

105-106;

140pp; English.

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Alignment
Pred. No.:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cystadencoarcinoma, endometrioid carcinoma, undifferentiated carcinoma, clear cell adenocarcinoma, cystadenofibroma, adenofibroma and Brenner tumour. The ovarian tumour marker genes of the invention were identified using SAGE (serial analysis of gene expression) and were found to be overexpressed in a broad variety of ovarian epithelial tumour cells relative to normal ovarian epithelial cells. The marker genes are implicated in immune response pathways, in the regulation of cell proliferation and in protein folding, and many of these are membrane-localised or secreted. In addition to their use as diagnostic and prognostic markers, the ovarian tumour marker genes or their encoded proteins may be used as therapeutic targets for the treatment and prevention of ovarian cancer. Sequences ABAB3081-ABAB3122, ABAB3180, ABAB3182 and ABAB3184 represent the ovarian tumour marker genes of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               at increased risk for developing ovarian cancer, in prognostic tests assessing the relative severity of ovarian cancer, in tests for monitoring a patient in remission from ovarian cancer and in tests for monitoring disease status in a patient being treated for ovarian cancer the methods can additionally be used to identify a particular tumour of the methods can additionally be used to identify a particular tumour over the methods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         being an ovarian tumour (i.e., an epithelial ovarian tumour selected from serous cystadenoma, borderline serous tumour, serous cystadenocarcinoma, mucinous cystadenoma, borderline mucinous tumour, serous cystadenocarcinoma,
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                                                       LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr
                                                                                                                                       GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
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                                                                                                                                                                               New apoE humanized mammalian useful for treating or prever arteriosclerosis -
                                                                                                        present sequence is the coding sequence for in the method of the present invention. The can be used for screening for agents useful Alzheimer's disease and arteriosclerosis.
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P-PSDB; AAB80997.
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                                                                                       Sequence 1156
                                                                                                                                            The present invention
                                                                                                                                                              Disclosure; Page 13-14;
                                                                                                                                                                                                                                                                                                                      JP2001017028-A.
                                                              No . :
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Conservative:
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Location/Qualifiers
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                                                                                                                                                                                                                          Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                       The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE3 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LI receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or ha defect in remnant removal.
                                                                                                                                                                                                                                                                                     Sequence 1156
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04-OCT-2000; 2000US-0679088
05-APR-2001; 2001US-0827854
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                               CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
                                                                     GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTTGGGATTACCTGCGCTGGGTGCAGACA
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(UYBO-)
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Alignment Scores:
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P-PSDB;
                                                                                      The present invention relates to an ApoE humanised mammalian cell. The present sequence is the coding sequence for human ApoE2, which was us in the method of the present invention. The ApoE humanised mammalian can be used for screening for agents useful for treating or preventin Alzheimer's disease and arteriosclerosis.
                                                                                                                                                      New apoE humanized mammalian useful for treating or preve arteriosclerosis -
                                                                       Sequence
                                                                                                                                       Disclosure;
                                                                                                                                                                                                                              06-MAY-1999;
                                                                                                                                                                                                                                             28-APR-2000;
                                                                                                                                                                                                                                                                            JP2001017028-A.
                                                                                                                                                                                                                                                                                                                                                   Human;
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DB; AAB80996.
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RESULT 8
AAD22049
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AC AAD2
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DT 12-F
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DE Huma
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KW Huma
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OS Homo
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                                                Human
                                                                                               AAD22049
         Homo
                      Human; apolipoprotein hypertriglyceridaemia;
                                                                12-FEB-2002
                                                                                AAD22049;
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                                                                                                                                                                                                                                                           LeuGluGluGlnAlaGlnGlnIleArgLeuGlnAlaGluAlaPheGlnAla
                                                                                                                                                                                                      LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
                                                                                                                                                                                                                                                                                                                                                                   ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                                               CTGGAGGAGCAGGCCCAGCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCC
                                                                                                                                                              AGCCGGACCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCCAAG
                                                                                                                                                                      SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
                                                                                                                                                                                              ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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                                                                                                                                                                                                                                                                                                                                                                                          ACCCCGGTGGCGGAGGAGACGCGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                  LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                                               apolipoprotein E (apoE)
                                                                                               standard;
                                                               (first
                                                               entry)
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                        E; apoE; cholesterol;
low density lipoprote
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                                                                                               ВP
                                              isoprotein, apoE2
                        lipoprotein;
                        atherosclerosis;
ein; LDL; ds.
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277

840 260 780 240

720 220 660 200 600 180 540 160 480 140

420 120 360 100 300 80 240 60 180 40

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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04-OCT-2000;
05-APR-2001;
                                                                                                                                                                                                                                                                          The present sequence is a human apolipoprotein E (apoE) isoprotein, apoE2 DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (I receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or hadefort in remnant removal
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 200
P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                          New apolipoprotein E polypeptide and nucleic acid, useful for l cholesterol, delaying the onset of or treating atherosclerosis mammal, without inducing hypertriglyceridemia -
                                                                                                                                                                                                                                                                                                                                                                                                                                                           (KOSP-)
(UYBO-)
                                                                                                                                                                                                                                                                                                                                                            Claim
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                  241
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                                                                                      GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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DB; AAE13295.
                          LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
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LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGlnLeu
                  CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
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2000US-0679088.
2001US-0827854.
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61..114
/*tag= b
115..1011
/*tag= c
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RESULT 9
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DE Hum
XX Hyp
XX Key
FT CDS
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FT mat.
FT mat.
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FT MAC2
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XX 06--
PR 06--
06-APR-2000;
04-OCT-2000;
05-APR-2001;
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                                              18-OCT-2001
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                                                                                           mat_peptide
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; 2001US-0827854.
                               2001WO-US11358
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is a human apolipoprotein E (apoE) allele, apoE2* DNA. The apoE lipoproteins are useful for lowering cholesterol, delaying the onset of atherosclerosis, treating or regressing atherosclerosis without inducing hypertriglyceridaemia, in a mammal lacking an endogenous, normally functioning apoE gene or low density lipoprotein (LI receptor or is at risk for developing atherosclerosis due to accumulation of lipoprotein remnants in the bloodstream or had defect in remnant removal
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(UYBO-)
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ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
                                                                                                         ThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAla
                                                                ACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCC
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haemolymph; lip
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DB; AAR86791.
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                                                                                                                     of recombinant apo:lipoprotein a sexta larvae with recombinant r polyhedrosis baculovirus vector
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                                                                                                 Columns 11-14; 10pp;
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Recombinant human apolipoprotein-E (ApoE) (AAR86791) can be produce by preparing a genetic construct (conty an ApoE-encoding sequence e.g. AA706957, and flanking regulatory sequences enabling the prot to be expressed in insect cells), which is then introduced into a Manduca sexta larva (using a recombinant Autographica californica nuclear polyhedrosis virus) and recovering the protein from the haemolymph of the larval host. The ApoE produced is in a form sufficiently complexed with lipids to be biologically active, whice cannot be achieved in insect cell cultures, and can therefore be

sequence, produced

protein

which be

in

applications

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RESULT 11
ABN95746
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                                                                  LeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMetGly
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CC The invention relates to a novel method for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma or metastatic liver cumour in a patient, and differentiating metastatic liver cancer from CC hepatocellular carcinoma in a patient, involving detecting the level of CC expression of two or more genes represented in ABN93503-ABN97455 in a CC tissue sample. The method of the invention has hepatotropic, and CC cytostatic activity. The method is useful for diagnosing and detecting CC the progression of liver cancer, hepatocellular carcinoma and metastatic CC liver carcinoma in a patient. The method is useful for identifying CC expression profiles which serve as useful diagnostic markers as well as CC markers that can be used to monitor disease states, disease progression, CC drug toxicity, drug efficacy and drug metabolism.

CC Note: The sequence data for this patent did not form part of the printed CC specification, but was obtained in electronic format directly from WIPO CC at ftp.wipo.int/pub/published_pct_sequences.
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disease progress:
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CC The method is based on changes in gene expression in BPH tissue isolated CC from patients exhibiting different clinical states of prostate CC hyperplasia as compared to normal prostate tissue. (I) comprises CC detecting the expression levels of one or more genes in prostate cells CC from the subject that are differentially regulated compared to normal prostate cells. (II) comprises preparing a first gene expression profile CC of BPH cells or BPH-like cell population, exposing the cells to the CC agent, preparing a second gene expression profile of the agent exposed CC cells, and comparing the first and second gene expression of BPH. (II) is CC useful for identifying an agent that modulates the onset or progression CC is useful for diagnosing the onset or progression of BPH. The methods are useful to present information identifying CC the expression level in a tissue or cells, by comparing the expression CC level of genes given in the specification in the tissue or cells to the compared to the expression of gene in the database, and displaying the compared to the expression levels of at least one gene in the tissue or cell sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         progression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying drugs for and diagnosing benign prostatic hyperplasia, by detecting expression levels of one or more genes in prostate cells from patient that are differentially regulated compared to normal prostate
                                                                       treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e invention relates to a method of diagnosing (I) the onset or or or screening identifying an agent that modulates the onset or progression
                                            to the expression level in BPH. Agents using (II) are useful BPH or prostate cancer. ABK64106-ABK64860 represent human rostatic hyperplasia gene sequences of the invention.
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1157 B₽; 212 Α; 370 Ç; 426 G; 149 H, 0 other;

prostatic

Pred. Alignment

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Best Local Similarity:
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stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous;
cytostatic; gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;
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Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature gene set

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Claim 1; SEQ ID 3787; 44pp; English.

The present invention describes a method (M1) for screening for an anti-neoplastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change in expression of at least one gene (I) of a signature gene set, where (I) comprises a sequence (S) selected from 8447 sequences (given in ABL6164 to ABL70110), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has cytostatic activity and can be used in gene therapy. M1 can be used for screening an anti-neoplastic agent, and can be used for producting a product which is the data collected with respect to the anti-neoplastic agent as a result of M1, and the data is sufficient to convey the chemical creatment of cancer such as colon, breast, stomach, lung, thyroid, oesophageal, ovarian, kidney, prostate or pancreatic cancer, infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.

Sequence 1157 BP; 212 A; 370 c; 426 <u>ن</u> 149 Τ, 0 other;

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Alignment Scores:
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                                                                                                                                                              The present invention relates to an ApoE humanised mammalian cell. The present sequence is the coding sequence for human ApoE4, which was us in the method of the present invention. The ApoE humanised mammalian can be used for screening for agents useful for treating or preventin Alzheimer's disease and arteriosclerosis.
                                                                                                                                                Sequence 1156 BP;
                                                                                                                                                                                                                              New apoE humanized mammalian cell useful for screening useful for treating or preventing Alzheimer's disease arteriosclerosis -
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                                                                                                                                                                                                              Disclosure; Page 16-17; 22pp;
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Search completed: March 14, 2003, 12:43:02 Job time : 161.176 secs

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Title:
Perfect score:
Sequence:
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MODEL-frame+_p2n.model -DEV-xlp

-MODEL-frame+_p2n.model -DEV-xlp

-O-/cgn2_1/USPTO_Spool_US09827854/runat_11032003_101611_27510/app_guery.fasta_1.3576

-DB=Issued_Patents_NA -OPMT=fastap | SUFFIX=P2n.rni -MINMATCH=0.1 -LOOPCL=0

-LOOPEXT=0 -UNITS-bits -START=1 -EMD=-1 -MATRIX=blosum62 -TRANS-human40.cdi

-LIST-45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THRIN=0 -ALIGN=15

-MODE-LOCAL -OUTFWI=pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000

-USER-US09827854 @CGN_1_1_224 @runat_11032003_101611_27510 -NCPU=6 -ICPU=3

-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120

-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=7

-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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INFORMATION FOR SEQ
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STREET: P.
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STATE: WI
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## ALIGNMENTS

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Sequence 1, Application US/07709949
Patent No. 5472858
GENERAL INFORMATION:
APPLICANT: Attie, Alan D
APPLICANT: Gretch, Daniel G
APPLICANT: Gretch, Daniel G
APPLICANT: Sturiey, Stephen L
APPLICANT: Beckage, Nancy E
TITLE OF INVENTION: Production of Recombinant Proteins in TITLE OF INVENTION: Insect Larvae
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSE: Quarles & Brady
STREET: P.O. Box 2113
CITY: Madison
STATE: WI
COUNTRY: USA
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: FORM:
MEDIUM TYPE: FORM:
MEDIUM TYPE: FORMS: PC-DOS/MS-DOS
SOFTWARRE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/709,949
FILING DATE: 1991064
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 9629691801
TELEPHONE: (608) 251-5000
TELEPAX: (608) 251-9166
INFORMATION FOR SEQ ID NO: 1:
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SEQUENCE CHARACTERISTICS:
LENGTH: 1157 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: double

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AUTHORS: McLean, J W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE
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DATE: 19:
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                 Score:
                             Pred. No.:
Percent Similarity:
                                           Alignment Scores:
                                                                           US-08-949-155-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                         TELEFAX: (713) 789-267
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: TA
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 61
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,
                                                                                                                                                                                                                                                                                                                                                            FILING DATE: Concurrently Herewitl CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 60/027,338
FILING DATE: 11-OCT-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Compositions and Methods for TITLE OF INVENTION: Generation of Transgenic Anim NUMBER OF SEQUENCES: 51
                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Piedrahita, Jora APPLICANT: Bazer, Fuller W
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CITY: Houston
STATE: TX
COUNTRY: US
ZIP: 77210-4433
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                                                                                       NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                  TYPE:
                                                                                                                                   TOPOLOGY:
                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                        TELEPHONE:
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                                                                                                                                                                               LENGTH: 1126 base pairs
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(713) 789-2679
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              Length:
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Version #1.30

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RESULT 3
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                                                                     Sequence 5, Application US/09819964
Patent No. 6369294
GENERAL INFORMATION:
APPLICANT: Piedrahita, Jorge
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                   APPLICANI. FILLER W Bazer, Fuller W TITLE OF INVENTION: Compositions and Methods Generation of Transgenic
CORRESPONDENCE ADDRESS:
              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                         GluMetGlySerArgThrArgAspAfgLeuAspGluValLysGluGlnValAlaGluVal 257
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72.96%
              Generation
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Percent Similarity:
Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alignment Scores:
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APPLICATION NUMBER: 08/949,155
FILING DATE: <UNKNOWN>
APPLICATION NUMBER: US 60/046,094
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                  GluGlnAlaValGluThrGluProGluProGluLeuArg-----GlnGlnThrGlu 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: TAMK:177
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                          GluGlnLeuThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAla 117
                                                                                                                                                ValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGlu 77
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              GCGCAGCTGGGCCCCGTGACCCAGGAGACGCAGGCGCCCTGTCCAAGGAGCTGCAGGCG
                                                                                  LeuArgAlaLeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGlu 97
                                                                                                                                                                                                            TrpGlnSerGlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrp
                                                                                                                                  GTGCAGTCCCTGTCTGACCAAGTGCAGGAGGAGCTGCTCAGCACCAAGGTCACCCAGGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/819,964 FILING DATE: 28-Mar-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: P.O. Box 4433 CITY: Houston
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COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
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Length: Matches: Conservative: Mismatches: Indels:

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GENERAL INFORMATION:
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                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: 1 Financial Center
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 0211
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPER: US/08/726,306A
FILING DATE: 02-Oct-1996
PRIOR APPLICATION NUMBER: US/08/726,306A
FILING DATE: 02-Oct-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
FILING DATE: 02-Oct-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/009,832
FILING DATE: 01-Jan-1996
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 96,048-A (3)
TELECOMMUNICATION INFORMATION:
                                      TELEFAX: (617) 345-911 INFORMATION FOR SEQ ID NO:
                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: van Leeuwen, Frederik Willem
APPLICANT: Burbach, Johannes Peter Henri
APPLICANT: Grosveld, Franklin G.
TITLE OF INVENTION: DIAGNOSIS METHOD AND REAGENTS
NUMBER OF SEQUENCES: 189
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5. 5958684
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517) 345-9111
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                                                                                                                                              Sequence 51, Application US/089
Patent NO. 6271436
GENERAL INFORMATION:
APPLICANT: Piedrahita, Jorg
APPLICANT: Bazer, Fuller W
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STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
STATE: TA
COUNTRY: US
COUNTRY: T7210-4433
                                                                                                        TITLE OF INVENTION: Compositions and Methods for TITLE OF INVENTION: Generation of Transgenic Anim NUMBER OF SEQUENCES: 51
                                                                                            CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                         LysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAlaValTyr 180
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                                                   Houston
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EDNESS: double
                                                                  P.O. Box 4433
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Indels:
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120

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241 60 181

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Animal

Species

199

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78		Qy 78
2896	TTGGGGGGAAAAAACTTTTTTTTTTTTTTTTCATTTGACTTCATGTCTTGCTTTCTTT	Db 2837 TTGGGGGGAAAAACTTTTTC
78		Qy 78
2836	TCCTTGTCGCCAACTCTTGGGGGTCTGGGTCTCTGTTTCTTTTTTTT	Db 2777 TCCTTGTCGCCAACTCTTGGG
78		Qy 78
2776	CCCTCCTGGCGAACCGTGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG	Db 2717 CCCTCCTGGCGAACCGTGTGT
78		Qy 78
2716	3ACTCCCGCGCGCGCGCGCGCGCGCGCGCGCGCGCTGA	Db 2658 CT-GACGTAAGTGCCCACCCGACTC
78		Qy 78 Leu
77 2657	ValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGlu        :::      :::	Oy 58 ValGlnThrLeuSerGluGlnValG        :::      :::          Db 2598 GTGCAGTCCCTGTCTGACCAAGTGC
2597	TGGCAGGGCAGCCAGCCCTGGGAGCAGCCCTGGGCCGCTTCTGGGATTACCTGCGCTGG	Db 2538 TGGCAGGCAGCCAGCCCTGG
57	GGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrp	Qy 38 TrpGlnSerGlyGlnArgTrp
37 2537	IProGluProGluLeuArgGlnGlnThrGlu 	Qy 21 GluGlnAlaValGluThrGluProGluProGluLeuAr
20 2489	MetLysValLeuTrpAlaAlaLeuLeuValThrPheLeuAlaGlyCysGlnAlaLysVal	Qy 1 MetLysValLeuTrpAlaAla              Db 2448 ATGGCTGTGAACTGGCTCGGT
	-277) x US-08-949-155-51 (1-4267)	US-09-827-854-15_COPY_1_277 (1-
	Length: 4267 Matches: 192 Conservative: 35 Mismatches: 40 Indels: 139 Gaps: 4	Alignment Scores:  Pred. No.:  Scores:  Scores:  Scores:  Scores:  Scores:  Scores:  866.50  Percent Similarity:  56.05%  Best Local Similarity:  Query Match:  B:  4
·	disk patible DOS/MS-DOS (elease #1.0, version #1.30 A: US/08/949,155 US/08/949,155 US 60/027,338 1996 US 60/046,094 1997 US 60/046,094 1997 HALOTI 41,071 41,071 MATION: -3000 6799 178	COMPUTER READABLE FORM:  MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible COMPUTER: IBM PC compatible COMPUTER: Patentin Release COMPUTER: Patentin Release CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/94 FILING DATE: CONCURRENT APPLICATION UMBER: US 60/02 FILING DATE: 11-OCT-1996 PRIOR APPLICATION UMBER: US 60/04 APPLICATION UMBER: US 60/04 FILING DATE: 10-MAY-1997 APPLICATION UMBER: US 60/04 FILING DATE: 09-MAY-1997 ATTORNEY/AGENT INFORMATION: NAME: Hibler, David W. REGISTRATION UMBER: 41,071 REFERENCE/DOCKET NUMBER: TAM TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 418-3000 TELEFAX: (713) 789-2679 INFORMATION FOR SEQ ID NO: 51: SEQUENCE CHARACTERISTICS: LENGTH: 4267 base pairs TYPE: nucleac acid STRANDEDNESS: Single TOPOLOGY: linear

COUNTRY: US ZIP: 77210-4433 ; COMPUTER READABLE FORM: ; MEDIUM TYPE: Floppy disk ; COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30	
CORRESPONDENCE ADDRES ADDRESSEE: ARNOL STREET: P.O. Box CITY: Houston STATE: TX	., ., ., ., .,
TITLE OF INVENTION: Compo	
RESULT 6 15(-964-51) Sequence 51, App Patent No. 63692 GENERAL INFORM APPLICANT	E S C C C C C C C C C C C C C C C C C C
Qy 273 GluAlaPheGlnAla 277	
:::          3557 CAGCTGGAGGAGGTGCG	
253 GlnValAlaGluValArgAlaLysLeuGluGluGlnAlaGlnGlnIleArg	Qy
Qy 233 ArgAlaArgMetGluGluMetGlySerArgThrArgAspArgLeuAspGluValLysGlu 252            :::	
Oy 213 ValGlySerLeuAlaGlyGlnProLeuGlnGluArgAlaGlnAlaTrpGlyGluArgLeu 232 ::: :::              :::          :::     Db 3437 CTGAGTACCAGGGCCGGCCAGCCGCTGCGCGCAGCGCGGAAGCCTGGGGCCAGAAGCTG 3496	
Qy 193 SerAlaIleArgGluArgLeuGlyProLeuValGluGlnGlyArgValArgAlaAlaThr 212	
Qy 173 LeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeu 192	
Qy 153 ValArgLeuAlaSerHisLeuArgLysLeuArgLysArgLeuLeuArgAspAlaAspAsp 172	
Qy 133 LeuValGlnTyrArgGlyGluValGlnAlaMetLeuGlyGlnSerThrGluGluLeuArg 152	
Qy 113 LysGluLeuGlnAlaAlaGlnAlaArgLeuGlyAlaAspMetGluAspValCysGlyArg 132	
Qy 93 LysSerGluLeuGluGluGluEnThrProValAlaGluGluThrArgAlaArgLeuSer 112 :::                             :::	
7 CCCGCCCTCTCTGCGCCCAGGGAGCTGATAGAGGAGGCATGAAGGAGGTGAAGGCCTAC	
2957 AATCTCTGGCACGTCCTGGCCATCGCCAGCTCAGGAGCCCTCCTTCTCCCCCTCACCGCC 3	Db
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REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK:177
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/819,964
FILING DATE: 28-Mar-2001
CLASSIFICATION: <Unknown>
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LENGTH: 4267 base pairs
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                         ATCTTGAGCTCCTGCCTTCGCCTGTCTCTGGGTCAGTCTTGCCGTCCCTTGCTGTCTCTG
                                                                             TTGGGGGGAAAAACTTTTTCTTTTTCTTTTCATTTGACTTCATGTCTTGCTTTCCTTTCC
                                                                                                                                 CCCTCCTGGCGAACCGTGTGTTCTGGACCCTCAGGCTCCACCCGTCCGGGTTTCCTTCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
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                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: HORNES, Erik
APPLICANT: UHLEN, Mathias
                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                        TITLE OF INVENTION: CINUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
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ATTORNEY/AGENT INFORMATION:
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CITY: Alexandria
                          APPLICATION NUMBER: UPFILING DATE: 19920519
              CLASSIFICATION:
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Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
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US-07-849-389-6
                                                                                                                                                                                      Sequence 24, Application US/08617256 Patent No. 6043031 GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 330 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: Single
              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatik
                                                                                                                   APPLICANT: Kvster, Hubert
TITLE OF INVENTION: DNA DIAGNOSTICS
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEB: LAHIVE & COCKFIELD
ADDRESSEB: LAHIVE & COCKFIELD
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ORIGINAL SOURCE:
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TELECOMMUNICATION INFORMATION
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                                                                             CITY: Boston
STATE: Massachusetts
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ZIP: 02109-1875
                                                                                                         STREET:
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  OPERATING SYSTEM:
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TELEFAX: (703)683-4109
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                                                                                                                                                                                                                                                                                                                                                                                                                                 GlyAlaGluArgGlyLeuSerAlaIleArg 196
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compatible
PC-DOS/MS-DOS
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alignment Scores:
                                                                                                                                                                                                   Sequence 24, Application US/09287141 Patent No. 6197498
                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617)227-5941 INFORMATION FOR SEQ ID NO:
         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: March 17, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth A.
REGISTRATION UMBER: 35,430
REFERENCE/POOCET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: SQ
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                            APPLICANT: K ster, I
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                      ADDRESSEE: Heller E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
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                                                  COUNTRY: USA
ZIP: 92037-9103
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STRANDEDNESS: sing
TOPOLOGY: linear
OPERATING SYSTEM:
                                                                              STATE:
                                                                                            CITY:
                                                                                                       STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGGCCCGAGGGCCGCC
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                                                                             La Jolla
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4250 Executive Square,
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Spectrometry

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COUNTRY:

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                                                                                                                               Sequence 24, Application US/09431613
Patent No. 6221601
GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                              APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA D: NUMBER OF SEQUENCES: 33 CORRESPONDENCE ADDRESS:
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REFERENCE/DOCKET NUMBER: 247
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
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FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
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STREET: 4250 E:
CITY: La Jolla
STATE: CA
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STRANDEDNESS: single
TOPOLOGY: linear
                                                ADDRESSEE:
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                               Heller Ehrman White & McAuliffe 50 Executive Square, 7th Floor
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374.00
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Indels:
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Query Match:
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Sequence 24, Application US/09504245
Patent No. 6221605
GENERAL INFORMATION:
APPLICANT: K ster, Hubert
TITLE OF INVENTION: DNA Diagnost:
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INFORMATION FOR SEQ ID NO:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: ITEM COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002G
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
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SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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LENGTH: 252 base pairs
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ATTORNEY/AGENT INFORMATION:
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FILING DATE: 18-MAR-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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STRANDEDNESS: single
TOPOLOGY: linear
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REGISTRATION NUMBER: 33,77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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                                                                                                                           GAGCGCGGCCTC
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(617)227-5941
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NUMBER OF SEQUENCES:

GENERAL INFORMATION:

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US-09-287-682-24

: Sequence 24, Application US/09287682

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DB:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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STREET: 4250 Ex
CITY: La Jolla
STATE: CA
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 24736-2002J
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 2473
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELECHAMUNICATION INFORMATION:
TELEPHONE: 619-587-5360
TELECHAMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-5941
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INFORMATION FOR SEQ ID NO:
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LENGTH: 252 base pairs
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FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L
REGISTRATION NUMBER: 33,77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/287,682
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MEDIUM TYPE: Diskette
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STREET: """
CITY: La Jolla
""ATE: CA
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ADDRESSEE: Heller Ehrman White & McAuliffe
STREET: 4250 Executive Square, 7th Floor
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APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
TITLE OF INVENTION: DNA Diag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 18-MAR-PRIOR APPLICATION DATA:
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181
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                AspAlaAspAspLeuGlnLysArgLeuAlaValTyrGlnAlaGlyAlaArgGluGlyAla
                                                                                    GluGluLeuArgValArgLeuAlaSerHisLeuArgLysLeuArgLysArgLeuLeuArg
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GATGCCGATGACCTGCAGAAGTCCCTGGCAGTGTACCAGGCCGGGGGCCCGGGGGCGCCC
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US-09-287-679-24
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TELEFAX: 619-587-5360
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-5941
INFORMATION FOR SEO ID NO: 24:
SEQUENCE CHARACTERISTICS:
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APPLICANT: K ster, Hubert
APPLICANT: Little, Daniel P.
APPLICANT: Braun, Andreas
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie
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TITLE OF INVENTION: DNA Diagnostics Based on Mass Spectrometry
NUMBER OF SEQUENCES: 33
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                                                                                    109 AlaArgLeuSerLysGluLeuGlnAlaAlaGlnAlaArgLeuGlyAlaAspMetGluAsp 128
                129 ValCysGlyArgLeuValGlnTyrArgGlyGluValGlnAlaMetLeuGlyGlnSerThr 148
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CITY: I
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                                                                  GCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCCCGGCTGGGCGCGGACATGGAGGAC
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Conservative:
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                                                                                                                                           Alignment Scores: Pred. No.:
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                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 252 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-450-8400
TELEPAS: 619-587-5360
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/41
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                   TOPOLOGY: 1
MOLECULE TYPE:
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APPLICATION NUMBER:
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109 AlaArgLeuSerLysGluLeuGlnAlaAlaGlnAlaArgLeuGlyAlaAspMetGluAsp 128
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CITY: La.
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ILECOMMUNICATION
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REGISTRATION NUMBER:
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(617)227-5941
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Matches:
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Alignment Scores: 9.9e-29 Pred. No.: 9.9e-29 Score: 374.00 Percent Similarity: 97.62% Best Local Similarity: 97.62% Query Match: 27.22%	BER: 2 8400 60 60 7400 941 0: 24: CS: 115	LASSIII OR API PPLIC, ILING OR API PPLIC, ILING ILING ORNEY AME: AME:	COUNTRY: USA  ZIP: 92037-9103  COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: PAtentin Release #1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/287 FILING DATE:	RESULT 15 US-09-287-681-24 US-09-287-681-24 Sequence 24, Application US/0928768 Patent No. 6277573 GENERAL INFORMATION: APPLICANT: K ster, Hubert TITLE OF INVENTION: DNA Diagnos NUMBER OF SEQUENCES: 3 CORRESPONDENCE ADDRESS: ADDRESSEE: Heller Ehrman Whit. STREET: 4250 Executive Square CITY: La Jolla STATE: Ca	189	121 G 169 A 181 G	Db 2 GCACGGCTGTCCAAGGAGCTGCAGGCC  Qy 129 ValCysGlyArgLeuValGlnTyrArc
Length: 252 Matches: 82 Conservative: 0 Mismatches: 2 Indels: 1	36-2002F	99 66	1.0, Version #1.25 7,681	31 stics Based on Mass Spectrometry te & McAuliffe e, 7th Floor		0 00 44	GCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCCCGGCTGGGCGCGGACATGGAGGAC 61  ValCysGlyArgLeuValGlnTyrArgGlyGluValGlnAlaMetLeuGlyGlnSerThr 148

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-DB=-Published_Applications_NA -QFMT|-fastap_SUFFIX=p2n.rnpb -MINNATCH=0.1
-LOOPEL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRAMS=human40.cdi -LIST=45 -DOCALIGN=200 -FHR_SCORE=pct -FHR_MAX=100
-THR_MIN-0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=2000000000 -USER=US0927854 = CGN 1_1_298 = grunat_11032003_101612_27578
-NCPU=6 -LCPU=3 -NO_XLPXY -NO_MAN6 |-LARGEQUERY -NBG_SCORES=0 -MAIT -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
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Maximum Match 100%
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Sequence 129, App
Sequence 17, Appl
Sequence 8, Appli
Sequence 454, App
                                                                                                 Description
                                                                                               SOFTWARE: Patentin version 3.1
SEQ ID NO 129
LENGTH: 1156
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
FEATURE:
NAME/KEY: CDS
LOCATION: (61)...(1014)
OTHER INFORMATION:
US-09-870-759-129
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US-09-870-759-129
    Percent Similarity:
                                         Alignment Scores: Pred. No.:
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APPLICANT: TERMAN, David S
APPLICANT: TERMAN, David S
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT OF NEOPLASTIC DISEASE
FILE REFERENCE: 870759
CURRENT APPLICATION NUMBER: US/09/870,759
CURRENT FILING DATE: 2002-01-14
PRIOR APPLICATION NUMBER: US/09/870,759
CURRENT FILING DATE: 2000-05-30
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 166
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Sequence 17, Application US/09802640
Publication No. US20030036057A1
GENERAL INFORMATION:
APPLICANT: Braun, Andreas
APPLICANT: Bonsal Aruna
APPLICANT: Kleyn Patrick
TITLE OF INVENTION: GENES AND POLYMORI
TITLE OF INVENTION: CARDIOVASCULAR D
FILE REFERENCE: 24736-2048
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       CANT: Bonsal Aruna
CANT: Kleyn Patrick
OF INVENTION: GENES AND POLYMORPHISMS ASSOCIATED
OF INVENTION: CARDIOVASCULAR DISEASE AND THEIR (
                                                                           SerArgThrArgAspArgLeuAspGluValLysGluGlnValAlaGluValArgAlaLys
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; LENGTH: 1156
; TYPE: DNA
; ORGANISM: HOMO Sapien
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (61)...(1014)
; OTHER INFORMATION: Nucleotide Se
; OTHER INFORMATION: (APOE)
US-09-802-640-17
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CURRENT FILING DATE: 2001-03-09
NUMBER OF SEQ ID NOS: 122
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
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 CTACAGGAGCGGGCCCAGGCCTGGGGCGAGCGGCTGCGCGCGGGTGGGAGGAGATGGGC
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PRIOR FILLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILLING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
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TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without induci
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CORRENT FILING DATE: 2001-04-05
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; SEQ ID NO 454
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; TYPE: DNA
; ORGANISM: Homo sapiens
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Patent No. US200
                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: GENES DIFFERENTIALLY I FILE REFERENCE: PA-0028 US CURRENT APPLICATION NUMBER: US/10/044,090 CURRENT FILING DATE: 2002-01-09 NUMBER OF SEQ ID NOS: 850
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                                                                                                                                                                                                                                                                                                     OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                               NAME/KEY: unsure LOCATION: 46
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OTHER INFORMATION: Inc
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; SEQ ID NO 12
; LENGTH: 1156
; TYPE: DNA
; ORGANISM: HOMO :
US-09-827-854-12
                                                                                                           APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods fo
TITLE OF INVENTION: cholesterol levels wit
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
SEQ ID NOS: 19
SOFTWARE: FASTSEQ for Windows Version 4.0
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; Sequence 9, Application US/09827854
; Patent No. US20020123093A1
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods f
TITLE OF INVENTION: cholesterol levels wi
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
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PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows V.
SEQ ID NO 9
LENGTH: 1156
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APPLICANT: Zannis, Vassilis
APPLICANT: Kypreos, Kyriakos E.
APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: Coholesterol levels without inducing
FILE REFERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
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Best Local Similarity:
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CURRENT APPLICATION NUMBER: US/09/954,456
CURRENT FILING DATE: 2001-09-18
PRIOR APPLICATION NUMBER: US/60/233,617
PRIOR ETLING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: US/60/234,052
PRIOR APPLICATION NUMBER: US/60/234,923
PRIOR FILING DATE: 2000-09-25
PRIOR PELLING DATE: 2000-09-25
PRIOR PELLING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR FILING DATE: 2000-09-26
PRIOR PELING DATE: 2000-09-26
PRIOR APPLICATION NUMBER: US/60/235,711
PRIOR APPLICATION NUMBER: US/60/235,711
PRIOR APPLICATION NUMBER: US/60/235,720
PRIOR APPLICATION NUMBER: US/60/235,720
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US/60/235,840
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US/60/235,863
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US/60/235,863
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PRIOR APPLICATION NUMBER: US/60/235,863
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US/60/235,863
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: HOMO
US-09-954-456-760
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                                                                                       US-09-827-854-15_COPY_1_277 (1-277)
                                                                                                                                           Query Match:
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                                                                                                                                                                                                                                                                                                                  SEQ ID NO 760
LENGTH: 1157
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TITLE OF INVENTION: Sets
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NUMBER OF SEQ ID NO
SOFTWARE: Patentin
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                       ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAAGGCCAAGGTG
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APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEO ID NOS: 3950
SOFTWARE: PATENTIN Ver. 2.1
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LENGTH: 1157
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; TYPE: DNA; ORGANISM: Homo sapie; FEATURE; COTHER INFORMATION: GUS-09-880-107-2244
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Pred. No.:
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TITLE OF INVENTION: Compounds and methods fc
TITLE OF INVENTION: Condesterol levels wit
FILE REPERENCE: 07180/004003
CURRENT APPLICATION NUMBER: US/09/827,854
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER: DATE: 2000-04-06
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SOFTWARE: FastSEQ for
SEQ ID NO 7
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                                            CAGGCCATGCTCGGCCAGAGCACCGAGGAGCTGCGGGTGCGCCTCGCCTCCCACCTGCGC
                                                                                                 GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg
                                                                                                                                   CGGCTGGGCGCGGACATGGAGGACGTGCCGCCGGCCGGCGTGCAGTACCGCGGCGAGGTG
                                                                                                                                             ArgLeuGlyAlaAspMetGluAspValCysGlyArgLeuValGlnTyrArgGlyGluVal
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CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: US 09/679,088
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 09/544,386
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Kypreos, Kyriakos E.
TITLE OF INVENTION: Compounds and methods for lowering
TITLE OF INVENTION: cholesterol levels without induc.
FILE REFERENCE: 07180/004003
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ORGANISM: Homo
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GlyGlnArgTrpGluLeuAlaLeuGlyArgPheTrpAspTyrLeuArgTrpValGlnThr
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                                                                                          CTGTCTGAGCAGGAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
                                                                                                   LeuSerGluGlnValGlnGluGluLeuLeuSerSerGlnValThrGlnGluLeuArgAla
                                                                                                                                      GGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGGGATTACCTGCGCTGGGTGCAGACA
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Best Local Similarity:
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TITLE OF INVENTION: METHOD FOR GENETIC ANALY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 11926-022001
CURRENT APPLICATION NUMBER: US/09/967,013
CURRENT FILING DATE: 2000-10-25
PRIOR APPLICATION NUMBER: 60/206,613
PRIOR PILING DATE: 2000-05-23
NUMBER OF SEQ ID NOS: 91
SOFTWARE: FastSEQ for Windows Version 4.0
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LENGTH: 41907
TYPE: DNA
ORGANISM: Homo:
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{\tt TrpAspTyrLeuArgTrpValGlnThrLeuSerGluGlnValGlnGluGluLeuLeuSer}
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238	18 yGlnProLeuGlnGluArgAlaGlnÅlaTrpGlyGluArgLeuArgAlaArgMetGluGl	N	γQ
21515	CTGGGGCCCCTGGTGGAACAGGGCCGCGTGCGGGCCGCCACTGTGGGCTCCCTGGCC	214	Д
218	lyProLeuValGluGlnGlyÅrgValArgAlaAlaThrValGlySerLeuAlaG	ш	γQ
198 21455	78 aValTyrGlnAlaGlyAlaArgGluĠlyAlaGluArgGlyLeuSerAlaIleArgGluAr 	1 213	Db VQ
178 21395	58 SLeuArgLysLeuArgLysArgLeuLeuArgAspAlaAspAspLeuGlnLysArgLeuAl	1 213	Dp QA
158 21335	38 yGluValGlnAlaMetLeuGlyGln\$erThrGluGluLeuArgValArgLeuAlaSerHi 	212	P 6
138 21275	18 aGlnAlaArgLeuGlyAlaAspMetĠluAspValCysGlyArgLeuValGlnTyrArgGl 	212	g dq VQ
118 21215	98 uGlnLeuThrProValAlaGluGlu†hrArgAlaArgLeuSerLysGluLeuGlnAlaAl 	211	DP QA
98 21155	79 -ArgAlaLeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGl 	210	DP GA
21095	36 CCCCATCCCAGCCTTCTCCCCGCCTCCCACTGTGCGACACCCTCCCGCCCTCTCGGCCG	210	В
78	78		Qy
21035	76 GCCTCTGCCCCGTTCCTTCTCTCCCCTCTTGGGTCTCTCTGGCTCATCCCCATCTCGCCCG	209	Дb
78	78		Qy
20975	16 TCGTCTCTGCCTCTGCCTCTGCATCTGCTCTCTGCATCTGTCTCTGTCTCTCTC	209	Дb
78	78		Qγ
20915	56 CCTCCCAAAGTGCTGGGATTAGAGGCATGAGCCACCTTGCCCGGCCTCCTAGCTCCTTCT	208	DЬ
78	78		Q,
20855	96 CTCACTGTGTGCCCAGGCTGGTCTTGAACTTCTGGGCTCAAGCGATCCTCCCCCCCTCGG	207	В
78	78	•	δ
20795	36 CGTCCTGGCTCTGTCTGTCCTTCCCTAGCTCTTTTATATAGAGACAGAGAGAG	207	Дb
78	78	•	δÕ
20735	76 TGGAATTCTCTCTCAGCTTTGTCTCTCTCTCTCTCTCTCT	206	Db
78	78	•	δŌ
20675	16 CCTGGGTCTCTGCTGGTTCTAGCTTCCTTTCCCATTTCTGACTCCTGGCTTTAGCTCTC	206	Вb
78	78	•	δÖ
20615	56 GGCGGCTATACCTCCCCAGGTCCAGGTTTCATTCTGCCCCTGTCGCTAAGTCTTGGGGGG	205	ДЬ
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20495	36 TGGGATTACCTGCGCTGCAGTGCAGACACTGTCTGAGCAGGTGCAGGAGGAGGAGCTGCTCAGC	204	Дd

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LENGTH: 786
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CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1999-03-12
PRIOR FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 896 SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rosen et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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AlaGluAlaPheGlnAla
                                   GAGCAGGTGGCGGAGGTGCGCCAAGCTGGAGGAGCAGGCCCAGCAGATACGCCTGCAG
                                                GluGlnValAlaGluValArgAlaLysLeuGluGluGlnAlaGlnGlnIleArgLeuGln
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US-09-880-107-2491/c; Sequence 2491, Application US/09880107; Patent No. US20020142981A1; GENERAL INFORMATION:
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DB:
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Best Local Similarity:
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CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-38
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GENERAL INFORMATION:
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LENGTH: 478
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TITLE OF INVENTION: Cancer Gene Determination and
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
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ORGANISM: Homo sapiens
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                                                                                                             uThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAl 120
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PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2491
LENGTH: 478
TYPE: DNA
ORGANISM: Homo sapiens
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; NAME/KEY: unsure
; LOCATION: (1)..(478)
; OTHER INFORMATION: n = a or
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Query Match:
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TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
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APPLICANT: Vockle
APPLICANT: Scher:
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                                                       uThrProValAlaGluGluThrArgAlaArgLeuSerLysGluLeuGlnAlaAlaGlnAl 120
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                                            GACCCCGGTGGCGGAGGAGACGCGGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGAGN
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Scherf, Uwe
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Search completed: March Job time: 70.563 secs

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-MODEL-frame+_p2n.model -DBV=xlp
-Q-/cgn2_1/USPTO_spool/US09827854/runat_11032003_101610_27495/app_query.fasta_1.3576
-DB-EST -QFMT=fastap -SUFFIX-p2n.rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS-bits -START=1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN=15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER-US09827854_@CGN_1_1_8826_@runat_11032003_101610_27495 -NCPU-6 -ICPU-3
-NO_XUSP-US09827854_@CGN_1_1_8826_@runat_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THEAPSC1=1 -KAROPO+10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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16154066 seqs, 8097743376 residues
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Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being predicted by analysis of the total score distribution. printed,

## SUMMARIES

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## ALIGNMENTS

AUTHORS TITLE	REFERENCE	SOURCE ORGANISM	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BG763371	RESULT 1
NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)	Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 842)	human Homo sapiens	EST.	BG763371.1 GI:14074024	BG763371	mRNA sequence.	602735433F1 NIH_MGC_49 Homo sapiens cDNA clone IMAGE:4860585 5',	BG763371 842 bp mRNA linear EST 15-MAY-2001		

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GlnAlaMetLeuGlyGlnSerThrGluGluLeuArgValArgLeuAlaSerHisLeuArg 160
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Tissue Procurement: ATCC/DCTP/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium ()
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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High quality sequence stop: 817.
Location/Qualifiers
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Contact: Robert Strausberg, Ph.D.
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/lab_host="DHIOB (phage-resistant)"
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ECORI; cDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GCCACGAC(G). Size-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
II RT (Life Technologies). Note: this is a NIH_MGC
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/clone_lib="NIH_MGC_49"
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                                                                                                                                                                                                                                                             Email: cgapbs r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI)
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
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National Institutes of Health, M
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                        LeuMetAspGluThrMetLysGluLeuLysAlaTyrLysSerGluLeuGluGluGluLeu 100
                                                                                                                                                        CTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCCAGGAACTGAGGGCG
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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/db_xref="taxon:9606"
/clone="IMAGE:5312024"
/clone_lib="NIH_MGC_96"
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/lab_host="DH10B"
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Eukaryota; Metazoa; Chordata;
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 800)
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BM042094.1
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NIH-MGC http://mgc.nci.nih.gov/.
NATIONAL Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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/organism="Homo sapiens"
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/lab_bost="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: xhoI; Site_2:
EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor:
GCCACGAG(G). Library constructed by Ling Hong in the
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11762 row: a column: 11
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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National Institutes of Health, Mammalian
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Plate: LLAM11760 row: b column:
High quality sequence stop: 782.
Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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/clone="IMAGE:5300259"
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/lab_host="DH10B"
                                                                                                 /organism="Homo sapiens"
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Percent Similarity:
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                                                                            roLeuGlnGluArg-AlaGlnAlaTrpGlyGluArgLeuArgAlaArgMetGluGluMet
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                                                             CGCTACAGGAGCGGGCCCAGGCCTGGGGCGAGCGGATACGCGCGATGGACGAGCAGG
                                                                                                                                                                                       CAGGCCGGGGCCCGAGGGCGCGAGCGCGCTC-AGCGCCATCCGCGAGCGCCTGGGG
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BG774871
BG774871.1 GI:
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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602649975F1 NIH_MGC_40
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Clone distribution: MGC clone distribution information can
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//lab_host="DBIOB (phage-resistant)"
/note="organ: prostate; Vector: pOTB7; Site_1: XhoI;
/note="organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5 adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."

a 242 c 324 g 103 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4760827"
/clone_lib="NIH_MGC_40"
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                                                                                                                                                                                                                                           BI600563
BI600563.1
EST.
                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sl
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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                                    found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov plate: LLAM11726 row: g column: 10 High quality sequence stop: 783.
Location/Qualifiers
                                                                                                                                                        Contact: Robert Strausberg,
                                                                                                                                                                         NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5287329"
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                                    GlnAlaGlyAlaArgGluGlyAlaGluArgGlyLeuSerAlaIleArgGluArgLeu-Gl
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/lab_host="DH10B"
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       found through the I.M.A.G.E. Consortium/LLNL at: http://image_llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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602717942F1 NIH_MGC_49 Homo sapiens
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2: Cloned into EcoRI/XhoI sites using the following 5 adaptor: GGCAGGGGO; Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC
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/db_xref="taxon:9606"
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                                                                                                                                                                                                 mRNA sequence.
BG829472
BG829472.1 GI
                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 845)
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602763768F1 NIH_MGC_42
                                                                                 Contact: Robert Strausberg,
                                                                                          Unpublished (1999)
                                                                                                                                                                 Homo sapiens
cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
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ProLeuValGluGlnGlyArgValArgAlaAlaThrValGlySerLeuAlaGlyGlnPro
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI(XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MCG_Library. | "
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/clone_lib="NIH_MGC_42"
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/db_xref="taxon:9606"
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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602513830F1 NIH_MGC_16 Homo s
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: LLCM1419 row: k column: 08
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/db_xref="taxon:9606"
/clone="IMAGE:4645759"
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                                              Eukaryota; Metazoa; Chordata; Craniata; v
Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 790)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
                                                                                                                      Homo
            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Pa
                                  Contact: Robert Strausberg,
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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/clone_lib="NIH_MGC_96"
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                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI)
Toshiyuki and Piero Carninoi (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11691 row: 1 column: 04
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
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/clone_lib="NIH_MGC_95"
/tissue_type="hippocampus"
/lab_host="DH10B"
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Mammalla; Eutheria; Primates; Catarrhini;
1 (bases 1 to 757)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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High quality sequence stop: 757.
Location/Qualifiers
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CONA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NH_MGC Library."

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/clone="IMAGE:5557004"
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM10669 row: i column: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 907)
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National Institutes of Health, Mammalian
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/lab_host="DH10B"
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